Access DB#	
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SEARCH REQUEST FORM

Scientific and Technical Information Center

Requester's Full Na	ime:	Examiner # : Serial Number: Results Format Preferred (Date:
Mail Roy and Did-	Phone Number 30	Serial Number:	
	Room Location:	Serial Number: Results Format Preferred (c	ircle): PAPER DISK F-M
If more than one s	earch is submitted, please	prioritize searches in order o	of need.
Include the elected spec utility of the invention.	ies or structures lesson t	ms. acronyms, and registry numbers,	ne subject matter to be searched
Title of Invention:			
Inventors (please prov	ide full names):		
			· ·
Earliest Priority Fili	ng Date:		
For Sequence Searches appropriate serial number	Only Please include all pertinent info	rmation (parent, child, divisional, or iss	trad = many
spropriate serial number.	•		uea patent numbers) along with th
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TAFF USE ONLY	**************************************	**********	******
rcher: 0. Schve	ber NA Sequence (#)		where applicable
rcher Phone #: 272			
reher Location: Rems	en EUI AGI		
e Searcher Picked Up:	**	Questel/Orbit	
c Completed:	Litigation	Dr.Link	
rcher Prep & Review Time:		Lexis/Nexis	
rical Prep Time:	, unext	Sequence Systems Com	Inger_
	Patent Family	www/Internet	

PTO-1590 (8-01)



STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 119583

TO: Terra Gibbs

Location: rem/2d10/2c18

Art Unit: 1635

Mi

Friday, April 16, 2004

Case Serial Number: 09/954556

From: David Schreiber

Location: Biotech-Chem Library

Remsen E01A61 Phone: 272-2526

david.schreiber@uspto.gov

Search Notes		TCS:	A Committee Comm	
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STIC SEARCH RESULTS FEEDBACK FORM

Biotech-Chem Library

Questions about the scope or the results of the search? Contact the searcher or contact:

Mary Hale, Information Branch Supervisor Remsen Bldg. 01 D86 571-272-2507

Vol	untary Results Feedback Form
>	I am an examiner in Workgroup: Example: 1610
A	Relevant prior art found, search results used as follows:
	☐ 102 rejection
	☐ 103 rejection
	Cited as being of interest.
	☐ Helped examiner better understand the invention.
	☐ Helped examiner better understand the state of the art in their technology.
	Types of relevant prior art found:
	☐ Foreign Patent(s)
	Non-Patent Literature (journal articles, conference proceedings, new product announcements etc.)
>	Relevant prior art not found:
	Results verified the lack of relevant prior art (helped determine patentability).
	Results were not useful in determining patentability or understanding the invention.
Cor	nments:

Drop off or send completed forms to STIC-Biotech-Chem Library Remsen Bldg.



119583

Schreiber, David

From:

Gibbs, Terra

Sent:

Wednesday, March 31, 2004 4:20 PM

To:

Schreiber, David

Subject:

Sequence searcj request...

Hi David,

I have another request for a score over length search:

I need a length limited nucleotide sequence search against nucleobases 1479 through 1508 of SEQ ID NO:3 in USSN 09,954,556, where the returns are rank ordered based on the score over length/ratio as we've discussed. I need the lengths limited to hits between 8 and 50 nucleotides, and I'll take as many hits as you can import into excel (64,000?), and alignments for anything above .75 on the above ratio. Hope this is clear, please call me if it's not. I also need the interference databases searched if possible.

Terra Cotta Gibbs, Ph.D. Art Unit 1635 Remsen Building 2D10 571-272-0758

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$ Karn,J., Gait,M.J., Heaphy,S. and Dingwall,C.
$ Wiral (HIV) growth inhibition
Unclassified to the patent: US 6114109-A 17 05-SEP-2000;
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Karn, J., Gait, M. John., Heaphy, S. and Dingwall, C.
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                                                                                                                                                                        Unclassified.

1 (bases 1 to 18)

1 (bases 1 to 18)

Karn, J., Gait, M. John., Heaphy, S. and Dingwall, C. RNA oligonucleotides that bind HIV tat protein Patent: US :8821046-A 23 13-OCT-1998;

Location/Qualifiers
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AR048082
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Sequence
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15; Conserv
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                   AR108985
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1 (bases 1 to 17)
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/mol_type="unassigned DNA"
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1 (bases 1 to 18).

1 (bases 1 to 18).

Karn, J., Gait, M.J., Heaphy, S. and Dingwall, Overland (HIV) growth inhibition
Patent: US 6114109-A 23 05-SEP-2000;
Location/Qualifiers
                                                                VIRAL (HIV) GROWTH INHIBITION
PATENT: WO 9202228-A 17 20-FEB-1992;
Location/Qualifiers
                                                                                                                  synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 17)
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/mol_type="unassigned RNA"
/db_xref="taxon:32630"
                     /organism="synthetic construct"
/mol_type="unassigned RNA"
/db_xref="taxon:32630"
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Sequence 1
AR340497
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1 (bases 1 to 17)

Karn, J. and Prescott, C.D.

Methods and kits for discovery of RNA-binding Patent: US 6573045-A 1 03-JUN-2003;

Location/Qualifiers

1. .17
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Human immunodeficiency virus 1
Viruses; Retroid viruses; Retroviridae; Lentivirus; Primate
                                                                                                                                                                                                                                                                                                                         Unknown
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Methods and kits for discovery of RNA-binding Patent: JP 2002526032-A 1 20-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                        Unknown
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JP 2002526032-A/1
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/mol_type="genomic RNA"
/db_xref="taxon:11676"
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                                                                            'mol_type="unassigned RNA"
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88.2%; Pred. No. 11;
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RESULT 13
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AX325661
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                                                                                                                                                                                                                                                                                                    Solanum tuberosum (potato)
Solanum tuberosum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus
Human immunodeficiency virus
Viruses; Retroid viruses; Retroviridae; Lentivirus; Primate
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Location/Qualifiers
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Patent: WO 9964625-A 1 16-DEC-1999;
                                                                                                                                                                                                           single stranded oligonucleotides
Patent: WO 0192512-A 1799 06-DEC-2001;
                                                                                                                                                                                                                                   Targeted chromosomal genomic alterations in
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                                                                                                                                    /organism="Solanum tuberosum"
/mol_type="unassigned DNA"
/db_xref="taxon:4113"
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/mol_type="unassigned RNA"
/db_xref="taxon:12721"
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Oligoribonucleotide 18-mer.
A21030 A21030.1 GI:641332
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Targeted Ciromosomal genomic alterations
single stranded oligonucleotides
Patent: WO 0192512-A 1800 06-DEC-2001;
UNIVERSITY OF DELAWARE (US)
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AX325662.1 GI:18096421
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1 (bases 1 to 18)
Karn,J., Gait,M.John., Heaphy,S. and Dingwall,C.RNA oligonucleotides that bind HIV tat protein
                                                                                                                                   Sequence 13
AR048072
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Patent: WO 9202228-A 20 20-FEB-1992;
Location/Qualifiers
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synthetic construct
                                                      Unclassified.
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1 (bases 1 to 18)
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/mcl_type="unassigned RNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:4113"
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                       RS Karn, J. and Prescott, C.D.

Methods and kits for discovery of RNA-binding compounds

AL Patent: JP 2002526032-A 3 20-AUG-2002;

RIBOTARGETS LTD

OS HIV

PN JP 2002526032-A/3

PD 20-AUG-2002

PF 04-UN-1999 JP 2000553615

PF 05-JUN-1998 GB 9812196.5,02-MAR-1999 GB 9904790.4 PI

JONATHAN KARN, CATHERINE DENISE PRESCOTT

PC C1201/68,C12N15/09,G01N21/78,G01N33/53,G01N33/542,G01N33/566,

PC G01N37/00,

PC C12N15/00

CC Methods and kits for discovery of RNA-binding compounds FH

Key

Location/Qualifiers

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FT source 1. 18
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/mol_type="unassion
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/mol_type="unassigned DNA"
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Human immunodeficiency virus
Viruses; Retroid viruses; Rel
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Methods and kits for discovery of RNA-binding compounds
Patent: US 6573045-A 3 03-UN-2003;
Location/Qualifiers
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/mol_type="unassigned RNA"
/db_xref="taxon:12721"
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Nucleic acids and secreted polypeptides encoded thereby
Patent: WO 0110902-A 40 15-FEB-2001;
Curagen Corporation (US)
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Location/Qualifiers
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/mol_type="unassigned DNA"
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Karn, J., Gait, M. John., Heaphy, S. and Dingwall, C. RNA oligonucleotides that bind HIV tat protein Patent: US 5821046-A 19 13-OCT-1998;
Location/Qualifiers
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Karn, J., Gair, M. John., Heaphy, S. and Dingwall, C. RNA oligonucleotides that bind HIV tat protein Patent: US 5821046-A 21 13-OCT-1998;

Location/Qualifiers
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Karn,J., Gait,M.J., Heaphy,S. and Dingwall,C.
Wiral (HIV) growth inhibition
Viral (HIV) growth inhibition
Patent: US 6114109-A 18 05-SEP-2000;
Location/Qualifiers
Unknow...
Unknow...
Unclassified.
1 (bases 1 to 17)
1 (bases 1 to 17)
Karn,J., Gait,M.J., Heaphy,S. and Dingwall,C.
Viral (HIV) growth inhibition
Patent: US 6114109-A 19 05-SEP-2000;
Patent: US 6114109-A 19 05-SEP-2000;
Patent: Location/Qualifiers
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Enzymatic nucleic acid treatment to hepatitis C virus infection. BD209410
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Karn,J., Gait,M.J., Heaphy,S. and Dingwall,C.
Viral (HIV) growth inhibition
Patent: US 6114109-A 21 05-SEP-2000;
Location/Qualifiers
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1 (bases 1 to 14)

AUTHORS

Blatt,L., McSwiggen,J.A., Roberts,E., Pavco,P.A. and Mack Enzymatic nucleic acid treatment of diseases or condition to hepatitis C virus infection Patent: JP 2002512791-A 3000 08-MAY-2002;

RIBOZYME PHARMACEUTICALS INC

OS Hepatitis virus (hepatitis C virus)

PD 08-MAY-2002

PP 26-APR-1999 JP 2000545991

PR 27-APR-1999 US 60/083217,18-SEP-1998 US 60/100842

25-FEB-1999 US 60/257608,23-MAR-1999 US 09/274553 PI

LAWRENCE BLATT,JAMES A MCSWIGGEN,ELISABETH ROBERTS,PAMELA A P. PAVCO,

PI DENNIS MACEJAK

PC C12N9/00,A61K31/7105,A61K38/21,A61K40''

PC A61K37/66,

PC C12N15/00

CC Enzymar''
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/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Brysch, W. and Schlingensiepen, K.
ANTISENSE OLIGONUCLECTIDE PREPARATION METHOD PATENT: WO 9833904-A 1714 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
unclassified.

1 (bases 1 to 14)

Brysch, W. and Schlingensiepen, K.

AN ANTISENSE OLIGONUCLECTIDE PREPARATION METHOD
Patent: WO 9833904-A 1715 06-AUG-1998;
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A89566
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Apple,R.J., Erlich
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/mc1_type="unassigned DNA"
/db_xref="taxon:32644"
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JP 2001511000-A/1715.
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An antisense oligonucleotide
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BD067079.1 GI:22612682
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1 (bases 1 to 14)
Schlingensiepen, K.H. and Brysch, W.
An antisense oligonucleotide preparation method
Patent: JP 2001511000-A 1715 07-AUG-2001;
BICGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH
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Schlingensiepen,K.H. and Brysch,W.
Schlingensiepen,K.H. and Brysch,W.
An antisense oligonucleotide preparation method
Patent: JP 2001511000-A 1714 07-AUG-2001;
BIOGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH
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An antisense oligonucleotide preparation method Location/Qualifiers
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/mol_type="genomic_DNA"
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Matches 12; Conserv
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1 (bases 1 to 14)

Blatt, L., Mcswiggen, J.A., Roberts, E., Pavco, P.A. and Macejak, D. Enzymatic nucleic acid treatment of diseases or conditions relato hepatitis C virus infection
Patent: JP 2002512791-A 2984 08-MAY-2002;

RIBOZYME PHARMACEUTICALS INC
OS Hepatitis virus (hepatitis C virus)
PN JP 2002512791-A/2984
PD 08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                   PF 26-APR-1999 JP 2000545991
PR 27-APR-1998 US 60/083217,18-SEP-1998 US 60/100842 PR 25-PBB-1999 US 09/257608,23-MAR-1999 US 09/274553 PI LAWRENCE BLATT, JAMES A MCSWIGGEN, ELISABETH ROBERTS, PAMELA A PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            unidentified
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JP 2002512791-A/2984
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11arity 100.0%; Pred. No.
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07-AUG-2001
30-JAN-1998 JP 1998532533
31-JAN-1997 EP 97101531.8
KARL HERMANN SCHLINGENSIEPEN, WOLFGANG BRYSCH
                                                                                                                                                                                                              hepatitis C virus infection.
Key Location/One
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                                                                /organism="unidentified"
/mol_type="genomic RNA"
/db_xref="taxon:32644"
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                                                                                                                                      Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Score 10.8; D
Pred. No. 28;
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19-JUN-1998 US 60/08997,19-JUN-1998 US 60/090079 PR
19-JUN-1998 US 60/090035,19-JUN-1998 US 60/09093 PR
19-JUN-1998 US 60/08997,19-JUN-1998 US 60/090972 PR
19-JUN-1998 US 60/089978,19-JUN-1998 US 60/090942 PR
19-JUN-1998 US 60/08999,19-JUN-1998 US 60/090043 PR
19-JUN-1998 US 60/090942,19-JUN-1998 US 60/090036 PR
19-JUN-1998 US 60/090044,19-JUN-1998 US 60/090036 PR
19-JUN-1998 US 60/090044,19-JUN-1998 US 60/09038 PR
19-JUN-1998 US 60/090084,19-JUN-1998 US 60/09033 PR
19-JUN-1998 US 60/090080,19-JUN-1998 US 60/090077 PR
19-JUN-1998 US 60/090080,19-JUN-1998 US 60/090077 PR
19-JUN-1998 US 60/090078,19-JUN-1998 US 60/090047 PR
19-JUN-1998 US 60/090078,19-JUN-1998 US 60/090045 PR
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unidentified unidentified unclassified.
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G01N37/00,
PC C12N15/
CC Preparal
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1 (bases 1 to 10)

Roberts, B. L. and Shankara, S.

Preparation and use of superior vaccines

Patent: JP 2002534056-A 1327 15-OCT-2002;
                                                                                                             Sequence 72 from Pat
A71513
A71513.1 GI:4775125
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                                                                                                                                                                                                                                                                                                                                             AGAAGCCAGA 10
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JP 2002534056-A/1327
15-OCT-2002
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Key Location/Qualifiers
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19-JUN-1998 US 60/0900
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72 from Patent WO9813521.
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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AX623670/c
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AX003113
AX003113.1
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Fesce, R. and Consalez, G.
METHOD FOR THE DIFFERENTIAL SCREENING OF GENE EXPRESSION
PRIMED REVERSE TRANSCRIPTION-POLYMERASE CHAIN REACTION
Patent: WO 9813521-A 72 02-APR-1998;
FESCE RICCARDO (IT)
                                                                       Method for determining homeostasis of the skin Patent: WO 02053774-A 711 11-JUL-2002; Henkel Komo nandtygesellschaft auf Aktien (DE) Location/Qualifiers
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Location/Qualifiers
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synthetic construct
artificial sequences.
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                                                                                                                                                Petersohn, D. Conradt, M. and Hofmann, K.
                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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100.0%; Pred. No. 33;
tive 0; Mismatches
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AX629909
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AX629388/c
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AX629388
AX629388.1 GI:28457426
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                                                                                                                                                                                                                                                                     Homo sapiens (human)
                                                                                                                                                        Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 02053774-A 6950 11-JUL-2002; Henkel Kommanditgesellschaft auf Aktien (DE) Location/Qualifiers
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Patent: WO 02053774-A 6429 11-JUL-2002;
Henkel Kommanditgesellschaft auf Aktien (DE)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                    /organism="Homo sapiens"
/mol_type="unassigned DNA"
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/db_xref="taxon:9606"
                                                               31.3%;
90.9%;
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Pred. No. 3
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Pred. No. 39;
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RESULT 44

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AR167701/c
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AX631091/c
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AR167701.1
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Unclassified.
1 (bases 1 to 12)
Inoue, T.
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AX631091
AX631091.1 GI:28459135
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Sequence 7143 from Patent
AX630102
AX630102.1 GI:28458140
                                         Unknown
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                                                       Unknown.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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Pred. No. 39;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Method for amplifying DNA fragment, method for estimating state of microorganism existing and method for estimating state of microorganism existing and method for estimating state of waste Patent: JP 199276176-A 65 12-OCT-1999;
SANYO ELECTRIC CO LID, SOCIETY FOR TECHNO-INNOVATION OF AGRICULTURE FORESTRY AND FISHERIES
OS Unidentified
PN JP 199276176-A/65
PD 12-OCT-1999
PF 31-WAR-1998 JP 1998087652
PR KOICHI INOUE
PR KOICHI INOUE
C12015/09,B09B3/00,C12Q1/00,C12Q1/68,C12N15/00,B09B3/00 CC
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                                           unidentified
                                                          E38691.1 GI:18621353
JP 2000270867-A/65.
                                                                                         12 bp Method and device for amplifying E38691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           unidentified
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12 bp DNA linear PAT 18-JUN-2001
Method for amplifying DNA fragment, method for estimating state of
microorganism existing and method for estimating state of waste.
                             unidentified
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             unclassified.
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C12N15/09,B09B3/00,C12Q1/00,C12Q1/68,C12N15/00,B09B3/00
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                                                                                                                                                                                                                                                                                                                           /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Pred. No. 41;
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                                                                                                         DNA fragment
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E64117/c
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Best Local Similarity
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OS Unidentified
PN JP 2000270867-A/65
PD 03-OCT-2000
PF 19-MAR-1999 JP 1999
PR KOICHI INOUE
PC C12NL5/09,CL2M1/00,CC Strandedness: Singlific CC Topology: Linear;
FH Key Loc
FT Source /or
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                                                                                                                                                                                                                                                                                Method for amplifying DNA fragment, amplification apparatus of DNA fragment, method for assaying a group of microorganisms, method for analyzing a group of microorganisms, method for analyzing a group of microorganisms, and method for assaying contaminating substance Patent: JP 199341989-A 65 14-DEC-1999;

SANYO ELECTRIC CO LTD, SOCIETY FOR TECHNO-INNOVATION OF AGRICULTURE PORESTRY AND FISHERIES
OS Artificial Sequence
PN JP 199341989-A/65
PD 14-DEC-1999
14-DEC-1999 JP 1999069694
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12 bp DNA linear PAT 18-JUN-2001 Method for amplifying DNA fragment, amplification apparatus of DNA fragment, method for assaying a group of microorganisms, method for analyzing a group of microorganisms, and method for assaying
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1 (bases 1 to 12)
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C12N15/09,C12M1/00,C12Q1/68,C12N15/00
Strandedness: Single;
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C12N15/09,C12M1/00,C12Q1/68,C12N15/00
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                                                                                                                           /organism='Artificial
Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Location/Qualifiers
                                                        /organism="synthetic co
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Score 9.4; Di
Pred. No. 41;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Method for discriminating microorganisms, apparatus for discriminating microorganisms, method for preparing data base for discriminating microorganisms, and recording medium recorded with program for discriminating microorganisms Patent: JP 2001275700-A 10 09-OCT-2001; SANYO. BLECTRIC CO LTD, SOCIETY FOR TECHNO-INNOVATION OF AGRICULTURE FORESTRY AND FISHERIES OS Artificial Sequence PN JP 2001275700-A/10 PD 09-OCT-2001
PF 31-MAR-2000 JP 2000099482
PF 31-MAR-2000 JP 2000099482
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   discriminating microorganisms, microorganisms program and record medium for recording the sa Patent: WO 0175156-A 10 11-0CT-2001; SANYO ELECTRIC CO LTD, SOCIETY FOR TECHNO INNO
                                                 Method of discriminating microorganisms, apparatus for discriminating microorganisms, method of making database for discriminating microorganisms, microorganisms discriminating
                                                                                                                  synthetic construct
synthetic construct
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1 (bases 1 to 12)
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WO 0175156-A/10.
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/db_xref="taxon:32630"
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BD240229/c
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OS Artificial Sequence
PN W0 0175156-A/10
PD 11-0CT-2001
PF 27-MAR-2001 W0 2001JP002516
PF 31-MAR-2000 JP 00P 099482
PI TAKAKAZU INOUE
PC C12Q1/68,C12N15/10,G01N33/48,G01N27/
Primer
                                                                                                                                                               19-JUN-1998 US
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Roberts, B. L. and Shankara, S.

Preparation and use of superior vaccines
Patent; JP 2002534056-A 1647 15-OCT-2002;

GENZYME CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                 PF OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JP 2002534056-A/1647.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BD240229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mammalia; Eutheria; Primates;
                                                                                                  G01N37/00
                                                                                                                                   C12N1/19
                                                                                                                  C12N1/21, C12N5/10, G01N33/15, G01N33/50, G01N33/53, G01N33/566,
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27-MAR-2001 WO 2001JP002516
31-MAR-2000 JP 00P 099482
TAKAKAZU INOUE
C12Q1/68,C12N15/10,G01N33/48,G01N27/447,G06F17/30,C12M1/00
                                                   Preparation
Key
                                                                                                                                                BRUCE L ROBERTS, SRINIVAS SHANKARA C12N15/09, C12N15/09, A61K39/00, A61P35/00, A61P37/04, C12N1/15,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  JP 2002534056-A/1647
15-OCT-2002
                                     source
                                                                                   C12N15/00, C12N5/00, C12N15/00
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/organism='Homo sapiens
Location/Qualifiers
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/mol_type="genomic DNA"
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90.9%;
                                                                  and
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US 60/0900
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60/090078,19-JUN-1998
60/090076,19-JUN-1998
60/111715
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60/090080,19-JUN-1998
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60/090035,19-JUN-1998
60/089992,19-JUN-1998
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60/090041,19-JUN-1998 US
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60/090042,19-JUN-1998
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                                                                     use of
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                                                   use of superior vaccines 
Location/Qualifiers
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Catarrhini; Hominidae;
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Gaps

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Euteleostomi;

(human)'

PC 8 60/090079 60/089993 60/090072 60/089991 60/090048 60/090043

60/089853

60/090040

PR

60/089844 60/089833 60/090036

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REFERENCE
AUTHORS
TITLE
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BD248497
LOCUS
                                                      SOURCE
ORGANISM
                                                                                                           ACCESSION
                                                                                                                                                  RESULT 54
AR303300/c
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ORGANISM
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Best Local (
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                                                                                                                                                                                                                                    1493
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UNIVERSITY OF ROCHESTER

OS Artificial Sequence
PN JP 2002529082-A/11
PD 10-SEP-2002
PF 10-NOV-1998 JP 2000581183
PI MAURICE ZAUDERER
PC C12N15/09, A01K67/027, A61K35/76, A61K39
PC A61K39/395,
PC A61K39/395,
PC A61K39/395,
PC C12Q1/02,
PC G101N33/574, C12N15/00, C12N5/00
CC MR7
FH Key
FT Source

FT Source

Occanism='Artificial
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1 (bases 1 to 10)
Shimamoto, A., Furuichi, Y.,
Watahiki, M.
                                                                                                           AR303300
Sequence 25 from patent US
AR303300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            synthetic construct
synthetic construct
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                                                      Unknown.
                                                                     Unknown.
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                                         Unclassified.
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MAURICE ZAUDERER
C12N15/09,A01K67/027,A61K35/76,A61K39/00,A61K39/04,A61K39/12,
                                                                                                                                                                                                                                                               Conservative
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/mol_type="genomic DNA"
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                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                  /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                 GI:31692076
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100.0%; Pr
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3 6544736.
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               Shibata,Y.,
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o. 41;
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               Funaki,H.,
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               Ohara, E.
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AX510716
LOCUS
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AX471386/c
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KEYWORDS
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KEYWORDS
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Best Local S
Matches
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JOURNAL
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AX471386
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Sequence 4 from Patent WO0227027.
AX510716
                                                                                                                                                            Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                            N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Method for synthesizing cDNA from mRNA sample Patent: US 6544736-A 25 08-APR-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  synthetic construct
synthetic construct
Similarity 9; Conserv
                                                                                                           HENKEL KGAA (DE)
                                                                                                                   Hofmann,K., Conradt,M. and Petersohn,D Method for determining skin stress or Patent: WO 02053773-A 963 11-JUL-2002;
                                                                                                                                                                                                                              AX471386.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     artificial sequences.
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9; Conserv
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 Conservative 0;
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963 from Patent WO02053773.
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/mol_type="genomic |
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                                            /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                  /mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                        /note="Oligonucleotide primer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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100.0%; Pred. No.
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k; Score 9; DB 1
k; Pred. No. 44;
0; Mismatches
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DEFINITION
ACCESSION
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AX628452/c
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AUTHORS
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AX626985
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KEYWORDS
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AX625231/c
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Sequence 4026 from Patent
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/db_xref="taxon:9606"
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Sequence 9694 from Patent
AX632652
          Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 2053-774-A 9694 11-JUL-2002; Henkel Kommanditgesellschaft auf Aktien (DE)
                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                           AX632652.1 GI:28468267
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
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Amerson,H.V., Wilcox,P., Sederoff,R.R., O'Malley,D.M. and Grattapaglia,D. Methods for within family selection of deperennials using genetic markers Patent: US 5908978-A 20 01-JUN-1999;
Location/Qualifiers
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Keinath, A.P., Somai, B.M. and Dean, R.A.

Method of diagnosing gummy stem blight

chain reaction assay

Patent: US 6258537-A 6 10-JUL-2001;

Location/Qualifiers
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Sequence 6 from patent US
AR161933
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BD240233
LOCUS
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SOURCE
ORGANISM
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ACCESSION
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                     REFERENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria; Primates; Catarrhini; H

1 (bases 1 to 10)

Roberts, B.L. and Shankara, S.

Preparation and use of superior vaccines
Preparation and use of superior vaccines
Patent: JP 2002534056-A 1651 15-OCT-2002;

GENZYME CORP
OS Homo sapiens (human)
PN JP 2002534056-A/1651
PD 15-OCT-2002
PF 18-JUN-1999 JP 2000554749
PR 19-JUN-1998 US 60/090039,19-JUN-1998
PR 19-JUN-1998 US 60/090039,19-JUN-1998
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                                                                                        E54684.1 GI:22556167
JP 2001211883-A/36.
Homo sapiens (human)
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens (human)
              Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 10)
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JP 2002534056-A/1651.
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  Matsushima, K.,
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C12N1/21,C12N5/10,G01N33/15,G01N33/50,G01N33/53,G01N33/566,
                                                                                                                                                                                                                                                                                                                                               GAAGCCAGCC 10
                                                                                                                                                                                                                                                                                                                                                                                        GAAGCCAGAC 1497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BRUCE L ROBERTS, SRINIVAS SHANKARA C12N15/09, C12N15/09, A61K39/00, A61F35/00, A61F37/04, C12N1/15,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Preparation and use of superior vaccines Key Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C12N15/00, C12N5/00, C12N15/00
                                                                                                                                                                                                       normal liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /crganism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism='Homo sapiens (human)'.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Metazoa; Chordata; Craniata; Vertebrata; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and use of superior
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98 US 60/990039,19-JUN-1998 US 60/99041,19-JUN-1998 US 60/99097,19-JUN-1998 US 60/089997,19-JUN-1998 US 60/089992,19-JUN-1998 US 60/089992,19-JUN-1998 US 60/089991,19-JUN-1998 US 60/089090,19-JUN-1998 US 60/090042,19-JUN-1998 US 60/090042,19-JUN-1998 US 60/090044,19-JUN-1998 US 60/090044,19-JUN-1998 US 60/090044,19-JUN-1998 US 60/090044,19-JUN-1998 US 60/090078,19-JUN-1998 US 60/090078,19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-
  Hashimoto,S.,
                                                                                                                                                                                                          cell
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Kaneko, S.
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                                             Craniata; Vertebrata; Catarrhini; Hominidae;
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Yamashita, T
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60/080993
60/090072
60/090941
60/090043
60/090043
60/090834
60/089833
60/090077
60/090045
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                                                                       Euteleostomi;
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REFERENCE
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FEATURES
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AUTHORS
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ORGANISM
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ACCESSION
VERSION
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AR303303
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SOURCE
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ACCESSION
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AR181983/c
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JOURNAL
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JOURNAL
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Best Local (
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Best Local
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L Patent: JP 2001211883-A 36 07-AUG-2001;
SCIENCE & TECH AGENCY
OS Homo sapiens (human)
PN JP 2001211883-A/36
PD 07-AUG-2001
PF 31-JAN-2000 JP 2000023170
PF 31-JAN-2000 JP 2000023170
PI KOJI MATSUSHIMA, SHINICHI HASHIMOTO, SHUICHI KANEKO, TARO PI YAMASHITA
PC C12N15/09,C07K16/18,C12P21/02,C12N15/00
CC
FH Key Location/Qualifiers.
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                                        Method for synthesizing cDNA from mRNA sample Patent: US 6544736-A 28 08-APR-2003; Location/Qualifiers
                                                                                                                        1 (bases 1 to 10)
Shimamoto, A., Furuichi, Y., Shibata, Y.,
                                                                                                                                                                                         Unknown
                                                                                                                                                                                                                                                                              AR303303
Sequence 28 from patent
AR303303
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Similarity 90.0%;
9; Conservation
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10 (bases 1 to 10)

11 (bases 1 to 10)

12 (bases 1 to 10)

13 (bases 1 to 10)

14 (bases 1 to 10)

15 (bases 1 to 10)

16 (bases 1 to 10)

17 (bases 1 to 10)

18 (bases 1 to 10)

19 (bases 1 to 10)

19 (bases 1 to 10)

10 (bas
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                                                                                                                                                                       Unclassified.
                                                                                                                                                                                                                    Unknown.
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                                                                                                           Watahiki,M.
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  /organism="unknown'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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6544736.
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                                                                                                                             Funaki,H.,
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                                                                                                                               Ohara, E.
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AR3 04 485
LOCUS
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SOURCE
ORGANISM
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                      RESULT 70
AR382219
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AR303338/c
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Sequence
AR382219
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1496 ACTTCAGCAG 1505
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Bullerdiek, J., Van de Ven, W.J.M., Schoenmakers, H.F.P.M. and Mols, R. Multiple-tumor aberrant growth genes
Patent: US 6544784-A 110 08-APR-2003;
                                                                                                                                                                                                                                                                                                                                            Sequence 110 from patent AR304485
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 63 from patent US AR303338
                                                                                                                                                                                                                                                                                                                                AR304485.1 GI:31693633
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Shimamoto, A., Furuichi, Y., Shibata, Y.,
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Similarity 90.0%;
9; Conservation
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                                                                                                                                                                        /organism="unknown"
/mol_type="genomic DNA"
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AX152377
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AX152226/c
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Sequence 141 from Patent W00138577.
AXI52226
AXI52226.1 GI:14533877
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Keinath,A.P., Somai,B.M. and Dean,R.A.
Method of diagnosing gummy stem blight
chain reaction assay
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       Human transcriptomes
Patent: WO 0138577-A 292 31-MAY-2001;
The Johns Hopkins University (US)
Location/Qualifiers
                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                         Homo sapiens (human)
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Sequence 292 from Patent
AX152377
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                                                         Velculescu, V.E., Vogelstein, B. and Kinzler, K.W
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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AX153162/c
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KEYWORDS
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1484 CCAAGAAGCC 1493
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Sequence 1077 from Patent W00138577,
AX153162
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Patent: WO 0138577-A 1077 31-MAY-2001;
The Johns Hopkins University (US)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
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Saccharomyces cerevisiae
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycets;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.

1 (bases 1 to 10)
S Velculescu, V.E., Vogelstein, B. and Kinzler, K.W.
Characterization of the yeast transcriptome
Patent: JP 2001509017-A 53 10-JUL-2001;
THE JOHNS HOPKINS UNIVERSITY SCHOOL OF MEDICINE
S Saccharomyces cerevisiae (yeast)
PN JP 2001509017-A/53
PD 10-JUL-2001
PN JP 2001509017-A/53
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PD 10-JUL-2001
PN JP 201509017-A/53
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PF 22-JAN-1998 JP 1998532117
PR 23-JAN-1997 US 60/035917
PI VICTOR E VELCULESCU, BERT VOGELSTEIN, KENNETH W KINZLER PC C12N15/10, C12N15/31, C07K14/395, C12Q1/68, C12Q1/02 CC Characterization of the yeast transcriptome FH Key Location/Qualifiers
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Characterization of the yeast transcriptome.
BD065117
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Sequence 63 from Patent WO0212498.
AX377325
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                                                                                                                                                                                      /organism='Saccharomyces
Location/Qualifiers
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OS Homo sapiens (human)
PN JP 2002209591-A/633
PD 30-JUL-2002
PF 19-JAN-2001 JP 2001012328
PI KOJI MATSUSHIMA, SHINICHI HASH
                                                                                                                                                   VECTOR FOR THE EXPRESSION IN YEASTS OF INTERLEUKINE-2, YEASTS AND METHOD FOR PREPARING INTERLEUKINE-2 Patent: WO 8503723-A 10 29-AUG-1985; Location/Qualifiers
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BD167088.1 GI:27872900
JP 2002209591-A/633.
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KOJI MATSUSHIMA, SHINICHI HASHIMOTO, SHUICHI KANEKO, TARO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism='Homo sapiens
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                    /organism="unidentified"
/mol_type="unassigned DN
/db_xref="taxon:32644"
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Lemoine, Y., Sondermeyer, P., Loison, G., Aigle, M. and Lecocq, J.P.
Lemoine, Y., Sondermeyer, P., Loison, G., Aigle, M. and Lecocq, J.P.
Yeast-expression vectors for interleukin-2, transformed yeasts and
process for the preparation of interleukin-2
Patent: EP 0152358-A 9 21-AUG-1985;
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Wang,C.-G. and Hepburn,A.G.
Genetic sequence assay using DNA triple strand formation
Patent: US 5861244-A 342 19-JAN-1999;
Location/Qualifiers
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AR353840
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1 (bases 1 to Baric, R.S. and
                        Unclassified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="unidentified"
/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Directional assembly of large viral genomes and chromosomes Patent: US 6593111-A 15 15-JUL-2003;
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                                                                                                                                           Hofmann,K., Conradt,M. and Petersohn,D. Method for determining skin stress or skin Patent: WO 02053773-A 309 11-JUL-2002;
                                                                                                                                                                                              Eukaryola; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                            /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="genomic |
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/db_xref="taxon:9606"
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Sequence 16 from Patent EP1262565.
AX616455
AX616455.1 GI:28447498
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Homo sapiens
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/mol_type="unassigned DNA"
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/db_xref="taxon:9606"
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Sequence 3005 from Patent
AX625964
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Method for determining homeostasis of the skin
Patent: WO 02053774-A 2530 11-JUL-2002;
Henkel Kommanditgesellschaft auf Aktien (DE)
Location/Qualifiers
            Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 02053774-A 3005 11-JUL-2002;
                                                                                               Homo sapiens (human)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                 Eukaryota;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Henkel Kommanditgesellschaft auf Aktien
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/mol_type="unassigned DN
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                   Metazoa;
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Primates;
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                                                                  Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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AX626847/c
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Patent: WO 02053774-A 4134 11-JUL-2002;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                       Sequence 4695 from Patent AX627654
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            AX627654.1
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/mol_type="unassigned DNA"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 4967 from Patent
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                        Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 02053774-A 5549 11-JUL-2002, Henkel Kommanditgesellschaft auf Aktien (DE)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Sequence 6157 from Patent
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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AX629825
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Mammalia; Eutheria; Primates;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                   Homo sapiens (human)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                            Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 02053794-A 9054 11-UUL-2002; Henkel Kommanditgesellschaft auf Aktien (DE)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Best Local Similarity 90.0%; Pred. No. 54;

Matches 9; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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                                                                           misc_structure
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                                                                                                      trans-activation
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The invention relates to a method of determining if a compound binds to a target RNA by treating a test compound with a reporter (R) labelled with a donor or acceptor group and labelled target RNA, labelled with the complementary donor or acceptor group, and measuring the fluorescence from fluorescent groups associated with a compound:target RNA complex in presence of the test compound and comparing the result with a standard.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The HIV-1 TAR stem-loop sequence (see AAQ21425) was compared to that from HIV-2 (see AAQ21426). The only regions common to the two TAR structures are in the loop region and the U-rich bubble in the upper stem. This 17-mer was synthesised and can hybridies to a 14-mer (see AAQ24061) to mimic the known HIV-1 tat recognition sequence but without the apical loop. In an assay, the 17-mer plus 14-mer structure competed satisfactorily with full-length (59-mer) TAR for binding to tat. See AAQ21427-Q21435 for TAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New HIV growth inhibiting oligo:nucleotide(s) - comprising rna binding sequences capable of binding to tat protein within cells, and in assay to identify cpds. with tat binding.
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02-MAR-1999;
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11-APR-2000
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                                                                                                              ing compounds that bind infections, tumors and
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Best Local Similarity
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30-OCT-2000;
27-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chromosomal genomic alteration; genome altering oligonucleotide; PCR; sso-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin; primer amino acid over production; herbicide resistance; glyphosate resistance; imidazolinone herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; triazine resistance; disease resistance; porphyric herbicide resistance; triazine resistance; disease resistance; modified oil production; modified starch production; waxy starch; altered floral morphology; male-sterile plant; albino mutant; increased stearate production; reduced plantiate production; albino plant; here alterial production; reduced linolenic acid production;
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                                                                                                                  New oligonucleotides with moderneating plants with desired nutritional value, herbicide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Solanum tuberosum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK26439 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                       EB,
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                                                                                                                                                                                                                                                                                                    Gamper HB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              production genome altering oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                    DELAWARE
                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-0208538P.
2000US-0244989P.
2001US-00818875.
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                                                                                                                     modified nuclease-resistant termini, useful red phenotypes, e.g. stress tolerance, improvide or disease resistance, or modified oil
                                                                                                                                                                                                                                                                                                       3
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Pred. No. 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                               or modified oil
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Claim

7;

Page

151;

220pp;

English

Kmiec EB,

Gamper HB,

Rice MC,

Kim

4

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ABK26440/c
ID ABK264
XX
ABK264
XX
AC ABK264
XX
AC ABK264
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AC Chromo
KW O-meth
KW DNA re
KW DNA re
KW anino
KW inidaz
KW porphy
KW modifi
KW altere
KW increa
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Solanu
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OS O1-JUN
PR 01-JUN
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PR 30-OCT
PR 27-MAR
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FI Kmiec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC genetic sequence, which comprises a single-stranded oligonucleotide conving a DNA domain. The DNA domain has at least one mismatch with CC respect to the genetic sequence to be altered and further comprises CC chemical modifications of the oligonucleotide. The chemical modifications of the oligonucleotide. The chemical modifications consist of o-methyl modification, an LNA modification, two or more CC consist of these modification, an LNA modification, two or more CC directing repair or alteration of plant genetic information of any two or CC directing repair or alteration of plant genetic information. The CC oligonucleotides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abiotic stress tolerance, improved CC untritional value (e.g. altering amino acid content of plants or CC offerring amino acid over production), herbicide resistance (e.g. glyphosate resistance, imidazolinone and sulphonylurea herbicide resistance, modified oil production, modified starch production (e.g. increased starch or production of waxy starch), altered floral CC morphology (e.g. male-sterile plants) or modified starch production CC morphology (e.g. male-sterile plants) or modified fatty acid content of cell prints of plants or altering cliconucleotides are also useful for producing albino mutants for the allering oligonucleotides are also useful for producing albino mutants for the called the content of the content 
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                                                                                                  01-JUN-2000;
30-OCT-2000;
27-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chromosomal genomic alteration; genome altering oligonucleotide; PCR; ss. o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin; primer; abiotic stress tolerance; improved mutritional value; hygromycin; primer; amino acid over production; herbicide resistance; glyphosate resistance; imidazolinome herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; triazine resistance; disease resistance; porphyric herbicide resistance; modified oil production; modified starch production; waxy starch; altered floral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced palmitate production; albino plant; increased stearate production; reduced linolenic acid production; horteranteric arcaes
                                                                                                                                                                                                              01-JUN-2001;
                                                                                                                                                                                                                                                                  06-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                              Solanum tuberosum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Waxy starch production genome altering oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                               photosynthetic process.
                                                      (UYDE ) UNIV DELAWARE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                     2000US-0208538P.
2000US-0244989P.
2001US-00818875.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 4 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        46.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 13.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 No.
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genetic sequence, which comprises a single-stranded oligonucleotide having a DNA domain. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modifications consist of o-methyl modification, an LNA modification, two or more phosphorothicate linkages on a terminus, or a combination of any two or more of these modifications. The oligonucleotides are useful for directing repair or alteration of plant genetic information. The oligonucleotides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abiotic stress tolerance, improved nutritional value (e.g. altering amino acid content of plants or conferring amino acid over production), herbicide resistance (e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                disease resistance, modified oil production, modified starch production (e.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatty acid content (e.g. reduced palmitate, increased stearate or reduced linolenic acid). The oligonucleotides are also useful for producing albino mutants for the analysis of photosynthetic processes. This sequence represents a genome altering oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     conferring amino acid over production), herbicide resistance (e.g. glyphosate resistance, imidazolinone and sulphonylurea herbicide resistance, porphyric herbicide resistance or triazine resistance),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil
                                                                    05-JUN-1998;
02-MAR-1999;
                                                                                                                                                                                                                                                           Antiviral; antibacterial; antifungal; anticancer; detection; TAR; fluorescence resonance energy transfer; tat; HIV-1; Rev response e autoimmune disease; trans-activation regulatory region; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17
 Karn J,
                                                                                                                       04-JUN-1999;
                                                                                                                                                           16-DEC-1999.
                                                                                                                                                                                            WO9964625-A2
                                                                                                                                                                                                                            Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                HIV-1 TAR oligonucleotide target sequence
                                                                                                                                                                                                                                                                                                                                                                      15-SEP-2003
11-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ59072;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ59072 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an oligonucleotide for targeted alteration of
                                    (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1485 CAAGAAGCCAGACTTCA 1501
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2002-106307/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7; Page 151; 220pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17
                                    RIBOTARGETS LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    B₽;
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                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                                                  98GB-00012196.
99GB-00004790.
                                                                                                                       99WO-GB001761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                46.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 17;
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                                                                                                                                                                                                                                                               Rev response element; ion; ss.
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Example; Page 31; 82pp; English.
                                                                                                         Identifying compounds that bind to target RNA, potentially useful
                                                                                                     infections, tumors and
                                                                                                      autoimmune diseases.
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cc target RNA by treating a test compound with a reporter (R) labelled with a donor or acceptor group and labelled target RNA, labelled with the cc complementary donor or acceptor group, and measuring the fluorescence from fluorescent groups associated with a compound:target RNA complex in presence of the test compound and comparing the result with a standard. The oligonucleotide containing the HIV-1 trans-activation regulatory region (TAR) to which the HIV-1 Tat protein binds. The complex is labelled with c-carboxyfluorescein and is used as a target for the binding of a clabelled ADP-1 protein. Detection of the complex is by fluorescence resonance energy transfer (FRET). The method is used to identify compounds that interfere with interaction between the target RNA and cligands or proteins. Compounds that are identified are potentially useful for treating infections (viral, bacterial or fungal), cancer and autoimmune diseases. The compounds are preferably directed to the TAR and the region of the transfer of the TAR and the region of the transfer of the transfer to the TAR and the region of the transfer to the TAR and the region of the transfer of the TAR and the region of the transfer of the TAR and the region of the TAR and the transfer of the transfer of the TAR and The invention relates to replication. (Updated on 15-SEP-2003 to standardise a method of determining SS compound binds to r (R) labelled with

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Sequence 18 BP; 5 A; 4 C; 6 G; 0 T; 3 U; 0 Other;
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ВB 1;

Length 18

0

Query Match

밁 Ś Best Local Similarity
Matches 13; Conserv 1490 AGCCAGACTTCAGCAGC 1506 AGCCAGAUJUGAGCAGC 17 Conservative 46.0%; 76.5%; 2; Mismatches Score 13.8; Pred. No. 9 . . 2 Indels 0; Gaps

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RESULT 6
AAF74454
ID FA454
ID FA454
AAF7
AXC AAF7
XX AAF7
XX Hum
XX Hum
XX Hum
XX Hum
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X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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Human; PRO; PROX; cytostatic; immunomodulatory; reproduction; gene therapy; cell proliferation; differentiation disorder; c immune associated disorder; gestational disease; pre-clampsia PCR primer; sequencing primer; ss. pre-clampsia; cancer;

NO:40

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Homo sapiens
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WO200110902-A2
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15-FEB-2001.

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11-AUG-2000; 2000WO-US021857.
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11-AUG-1999; 99US-0148433P
10-AUG-2000; 2000US-00635949
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CURAGEN
CORP.
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RΑ,
Fernandes
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WPI; 2001-147509/15.
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Nucleic acids encoding secreted polypeptides, polypeptides, useful for treating a syndrome associated disorder, e.g. cancer.
                   , designated w
                                     PROX
                     ģ
                     PROX-
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Example

<u>ب</u>

Page 119; 166pp;

English

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RESULT 7
AAF74457/c
ID AAF74457 standard; D
XX
AAF74457;
AC AAF74457;
AC AAF74457;
XX
O9-MAY-2001 (first
XX
DE Human PRO2 gene-spec
XX
Human; PRO; PROX; cy
KW gene therapy; cell p
YM gene therapy; cell p
KW pCR primer; sequenci
OS Homo sapiens.
XX
PN WO200110902-A2.
XX
D15-FEB-2001.
XX
XX
11-AUG-2000; 2000W0-
XX
XX
11-AUG-2000; 2000W0-
XX
XX
11-AUG-2000; 2000W0-
XX
PF 11-AUG-2000; 2000W0-
XX
PF 10-AUG-2000; 2000W0-
XX
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PT polypeptides, useful
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CC The present inventio
CC polypeptides design
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                                                                                                                                                                                                                                Nucleic acids encoding secreted polypeptides, designated PROX polypeptides, useful for treating a syndrome associated with associated disorder, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene therapy; cell proliferation; differentiation disorder; ca immune associated disorder; gestational disease; pre-clampsia;
                             The present invention describes isolated nucleic acids encoding secreted polypeptides, designated PROX polypeptides (i.e. a PRO polypeptide where X is an integer from 1 to 17). PROX polypeptides have cytostatic, immunomodulatory and reproduction activities, and can be used in gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-AUG-1999; 99US-0148433P.
10-AUG-2000; 2000US-00635949.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PRO2 gene-specific sequencing primer SEQ ID NO:43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRO; PROX; cytostatic; immunomodulatory; reproduction;
herapy; cell proliferation; differentiation disorder; c
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                                                                                                                                                                             Page 119; 166pp; English
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88.2%;
     antagonists and PROX agonists.
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an be used in gene
PROX polypeptides,
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                                                                                                                                                                                                                                                  Sequences AAX57562-X57571 represent antisense oligonucleotides targeted to a region spanning 4-9 codons downstream of the AUG translation initiation codon of the human insulin-like growth factor-1 receptor (IGF-IR) gene. The antisense oligonucleotides inhibit the expression of IGF-IR, which in turn induces apoptosis, especially in a tumour cell. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX57566;
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                                                                                                                                                                     Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 16; 23pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human insulin-like growth factor-1 receptor gene antisense
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GAAGCCAGACTTCA 1501
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inhibition; induction; apoptosis; tumour; liposome;
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                                                                                                                                                                                                                                                                                                                              The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3], which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see ARP45151 and ARF45153-01igonucleotides of the present invention (see ARP45151) and ARF45153-1745 [The method is useful for ameliorating the effects of psoriasis, inthibyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, inthibyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, anyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                             Sequence
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GGAGCCAGACTTCA 2
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                                                                                                                                                                                                                                                 B₽;
                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Werther
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                                                                                                                                                                                                                                                                                                       other hyperplasia
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                                                                                                                                                                                                                                             4 C;
                                                                                                                                                41.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15
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                                                                                                                                                                                                                                                 G; 4 T;
                                                                                                                                             Score 12.4;
Pred. No. 1
                                                                                                                          Mismatches
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                                                                                                                                                       15;
                                                                                                                                                                                                                                                 0 Other;
                                                                                                                                                                                DB 1;
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                                                                                                                                                                             Length 15;
                                                                                                                          Indels
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                                                                                                                   Gaps
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RESULT 10 AAF49087/c

RESULT 11
AAZ64832/c
ID AAZ648
XX
AC AAZ648
XX

AAZ64832 standard; RNA; 14

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                                                                                                                    Best
                                                                                                Matches
                                                                                                                                        Query Match
                                                                                                                                                                                                                                           disease,
vessels c
                                                                                                                                                                                                                                                          skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see ARF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrheea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood careary active the retina, and sease, when the retina is a necessary of the skin, a capacity of the skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; hyperneovascular condition of the retina; ss.
                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-041421/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 8; Page 61; 201pp; English
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                                                                                                                      Local
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                                                                                             l Similarity
13; Conserv
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                                                GAAGCCAGACTTCA 1501
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  GGAGCCAGACTTCA
                                                                                                                                                                                                                                             any
                                                                                                                                                                                           BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention relates to a method for ameliorating the effects ers. The method comprises contacting the skin with an
                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Werther GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
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                                                                                                                                                                                           2 A;
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                                                                                                                    41.3%;
                                                                                                                                                                                           4 C;
                                                                                                                                                                                                                                           hyperplasia
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                                                                                                                                                                                           G; 4 T;
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                                                                                                                    Pred.
                                                                                                                         Score 12.4;
Pred. No. 1
                                                                                                  Mismatches
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                                                                                                                                             DB 1;
                                                                                                                                                                                                Other;
                                                                                                                                             Length 15;
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28-MAR-2000

(first

entry)

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RESULT 12
ABX01669/c
ID ABX016
XX ABX016
AC ABX016
XX 23-DEC
XX Hepati
XX Enzyma
KW HCV ri
KW Liver
KW Liver
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                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents the preferred target sequence of an enzymatic nucleic acid, especially a hairpin ribozyme, which cleaves the Hepatitis C virus (HCV) RNA sequence at the base position given in the descriptor line. The HCV sequence was screened for optimal ribozyme target sites using a computer folding algorithm and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised to target these sites and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes of the invention inhibit gene expression and/or viral replication, and are used to treat diseases associated with Hepatitis C virus (HCV) infection, e.g. cirrhosis, liver failure and hepatocellular carcinoma. The ribozymes may be used in combination with
              Enzymatic nucleic acid; RNA cleavage; Hepatitis C virus infection; HCV ribozyme; HCV expression; HCV replication; cirrhosis; virucide; liver failure; hepatocellular carcinoma; HCV infection; drug therap
                                                                                                                                                                                                                                                                                                                                                                                             hepatocellular carcinoma. The ribozymes may be used in combination with interferon to treat HCV infection, other infectious diseases, autoimmune diseases, and cancer
                                                                                                                                                                   ABX01669 standard;
                                                                                                                                                                                                                                                                                                                                                                   Sequence 14 BP; 3 A; 2 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel ribozymes for the treatment of hepatitis C infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Enzymatic nucleic acid; hammerhead ribozyme; virus replication; cleavag cirrhosis; liver failure; hepatocellular carcinoma; interferon; cancer; autoimmune disease; ss.
                                                                                                          23-DEC-2002
                                                                                                                                        ABX01669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; Page 99; 123pp; English.
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25-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RIBO-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for hairpin ribozyme which cleaves HCV at
                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                        (first
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99US-00257608.
99US-00274553.
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                                                                            substrate
                                                                                                                                                                     RNA;
                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                        40.0%;
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                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                          <u>.</u>
                                                                            #154 for HCV hairpin
                                                                                                                                                                                                                                                                                                                       Score 12; DB Pred. No. 16;
                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                       DB 1;
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                                                                                                                                                                                                                                                                                                                                       Length 14;
                                                                            ribozyme #154.
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 n; drug therapy; cytostatic;
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                                                                                                                                                                                                                                                                                                          Gaps
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WPI;
                                                                                                                                                                                                                                                                                         interferon gamma; consensus interferon; hepatotropic; antiinflammatory; substrate; hairpin ribozyme; HP ribozyme; ss.
                            Blatt L,
                                                                                                    (BLAT/)
                                                                                                                                               23-MAR-1999;
                                                                                                                                                                           23-MAR-1999;
                                                                                                                                                                                                        27-JUN-2002
                                                          (MACE/)
                                                                                        ROBE/)
 2002-617759/66
                                                                        ROBERTS B. PAVCO P A.
                                                                                                                    BLATT
                                                          MACEJACK D
                                                                                                    MCSWIGGEN J
                           Mcswiggen JA,
                                                                                                                                               99US-00274553.
                                                                                                                                                                           99US-00274553.
                              Roberts B,
                              Pavco PA,
                              Macejack D;
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enzymatic nucleic acid or ribozyme is in a hammerhead (HH) or hairpin (HP) motif where the binding arms comprise sequences complementary to one of the substrate sequences defined in the specification. The HCV ribozymes are useful for modulating the expression and/or replication of HCV. They can be used to treat cirrhosis, liver failure and/or hepatocellular carcinoma. The HCV ribozymes are also useful for treating a condition associated with HCV infection in conjunction with one or more interferon alpha, beta or gamma or consensus interferon. The present sequence represents a substrate for a HCV hairpin (HP) ribozyme. Note: Some of the sequence data for this patent did not form part of the printed specification. The complete sequence data for this patent was obtained in electronic format directly from the USPTO web site at New ribozymes targeting RNA derived from hepatitis C virus inhibit viral replication and are useful to treat hepatitis C virus infections and cirrhosis, liver failure or hepatocellular carcinoma. The present invention relates to enzymatic nucleic acids which specifically cleave RNA derived from Hepatitis C virus (HCV). Claim 2; Page 62; 80pp; English. (HCV). The

Sequence 14 BP; 3 A; 2 C; 6 G; 0 T; 3 U; 0 Other;

seqdata.uspto.gov/psipsDIDEntry.html

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В
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           Query Match
Best Local S
Matches 12
     1497
12
              Similarity
     CTTCAGCAGCCA 1508
CTTCAGCAGCCA 1
           ilarity 100.0%;
Conservative (
                 40.0%;
           ; Score 12; DB
; Pred. No. 16;
0; Mismatches
                 DB 1;
              16;
           0
                 Length 14;
           Indels
           ٥,
           Gaps
           0
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ARESULT 13
AAF49084/C
ID AAF490
XX AAF490
XX AAF490
XX AAF490
XX IGF-I
XX AAF490
KW CYCOST
KW GTOWTH
KW GTOWTH
KW KETATO
IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilar growth factor mediated cell proliferation; ichthyosis; serborrhoea; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disc
                                                                  Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis;
                                                                                                                                                                                                                                                     AAF49084;
                                                                                                                                                                                                                                                                                                   AAF49084 standard; DNA; 15
                                                                                                                                                                                                          30-MAR-2001
                                                                                                                                                             IGF-I oligonucleotide #44.
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pilaris;

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RESULT 14
AAF49085/c
ID AAF490
XX AAF490
XX AAF490
XX JO-MAR
XX ANTISE
KW CYTOST
KW SKIN d
KW SKIN d
KW IGP bi
KW GY-MAR
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XX HOP-
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Best Local :
                                                                                             Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; kelo skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pittyriasi IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris growth factor mediated cell proliferation; ichthyosis; serborrhoea; rukeratosis; neoplasia; seleroderma; wart; skin cancer; sclerotic diseas; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15
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                                                                                                                                                                                                                                                                                                                                                IGF-I oligonucleotide #45.
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                                                                      condition of the retina;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A;
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100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 G; 4 T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor, IGF binding protein [IGFBB]-2 or IGFBB3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see ARF45151 and ARF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia
                                                                                                                                      neutral lipid transfer; plasma lipoprotein; atherosclerosis; atherectomy; reverse cholesterol transport; high density lipoprotein; therapy; CETP; familial hypercholesterolaemia; dyslipidaemia; hyperalipolatecinaemia; peripheral vascular disease; hyperbetalipoproteinaemia; RCT; inhibitor; angioplastic restenceis; low density lipoprotein; diabetes; HDL; human;
                                                                                                                                                                                                                                                                                                   28-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                      AAT49619 standard; RNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           skin disorders. The method comprises contacting the skin with antisense oligonucleotide, (for Insulin-like Growth Factor [IC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 8; Page 61; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-041421/05
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                                                                                                                                                                                                                              Hammerhead ribozyme; cholesterol ester transfer protein; mRNA cleavage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MURD-) MURDOCH CHILDRENS RES
                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inflammation.
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12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15 BP; 4 A; 3 C; 4 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                 ribozyme target sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 effects of a disorder, e.g. psoriasis, by admir
treatment (optional) and an antisense nucleic
es growth factor mediated cell proliferation ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               40.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GA,
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                                                                                                                                                                                                                                                                                                                                                                        ВP
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 to a method for ameliorating the effects
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                                                                                                                                                                                                                                                                 #477.
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KW AFERS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC ester transfer protein (CETP) hammerhead (HH) ribozymes (see AAT49881-CC T50137). CETP is a 74 kD glycoprotein that facilitates neutral lipid CC transfer between plasma lipoproteins. The numbering of the targets refers to the position of the cleavage site in full length CETP. The ribozyme CC binds to 5 nucleotides either side of this site, provided the sequence UH is immediately upstream. The ribozymes are able to cleave mRNA from the CC gene encoding CETP, thereby blocking synthesis and/or expression of the CC gene encoding CETP, thereby blocking synthesis and/or expression of the CC gene inhibited (or eliminated) thereby preventing the reduction in size CC and therefore increasing HDL levels. The ribozymes can be used to treat conditions associated with abnormal levels of CETP, specifically familial hypercholesteroleemia, atherosclerosis, peripheral vascular disease, the peripheral vascular complications of diabetes, transplant, atherectomy and CC vascular complications of diabetes, transplant, atherectomy and consisty lipoproteins (LDL), and a corresponding increase in HDL levels. The levels of HDL lavels are favourably altered (a decrease in LDL levels, and a corresponding increase in HDL levels). The study genetic drift cartivity.

CC rectific activity
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Matches
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Best Local Similarity
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                                                                                                                    Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                       IGFBP3
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                                                                                                neovascular condition of the
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                                                                                                                                                                                                                                                                                                                                                                                       oligonucleotide #1259.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX57564;
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                                                                                                                 03-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-JUN-1999;
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                                                        04-NOV-1997;
                                                                                                                                                                         14-MAY-1999
                                                                                                                                                                                                                                  WO9923259-A1
                                                                                                                                                                                                                                                                                       Homo
                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                        expression;
                                                                                                                                                                                                                                                                                                                                                                                                        Antisense; human; insulin-like growth factor-1 receptor; IGF-1R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a method for ameliorating the effects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 7; Page 52; 201pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           oligo #3 to insulin-like growth factor I receptor
                                                                                                                                                                                                                                                                                                                                                                        inhibition; induction; apoptosis; tumour; liposome; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 5 A; 6 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000WO-AU000693
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                                                                                                                    98WO-US023418
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 11.8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                     isolated from Histoplasma capsulatum (HC). HC polypeptides can be used for detecting annibodies for detecting a previous or current HC infectin a subject They can also be injected into the skin of a subject to detect past exposure to HC by detecting swelling of the skin. The annibodies can be used for detecting current HC infection in a subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequences AAX57562-X57571 represent antisense oligonucleotides targeted to a region spanning 4-9 codons downstream of the AUG translation initiation codon of the human insulin-like growth factor-1 receptor (IGF-IR) gene. The antisense oligonucleotides inhibit the expression of IGF-
HC nucleic acids and polypeptides can also be used for the treatment of histoplasmosis as well as in vaccines for the prevention of histoplasmosis
                                                                                                                                                                                                                                                                                                                     New isolated Histoplasma capsulatum nucleic acids, used products for the diagnosis, prevention and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Histoplasma capsulatum; fungus; M antigen; vaccine; detection; histoplasmosis; diagnosis; infection; antimicrobial; antibody;
                                                                                                                                                                                                               The present sequence represents a degenerate PCR primer
                                                                                                                                                                                                                                                                      Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zancope-Oliveira RM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-313361/26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1488 GAAGCCAGACTTC 1500
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                                                                                                                                                                                                                                                                      Page 49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lott TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antigen degenerate PCR primer M4F.
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histoplasmosis.
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RESULT 19
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Best Local S
Matches 10
                                                                disease,
vessels (
                                                                                          oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
                                                                                                                                                                                                                                                                                                        skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP), which is capable inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonuclectide which can be used to design the antisense oligonuclectide which can be used to design the antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding procein; IGFBP2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cance; sclerotic disease; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-041421/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 8; Page 61; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UV (ultra-violet) treatment inhibits or reduces growth !
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hyperneovascular condition; hyperplasia; kidney disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       present invention relates to a method for ameliorating the effects
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Werther
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               reatment (optional) and an antisense nucleic acid growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       38.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Edmondson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  G; 2 T; 0 U; 5 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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Best Local Similarity
Matches 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                              The present sequence represents the preferred target sequence of an enzymatic nucleic acid, especially a hairpin ribozyme, which cleaves the Hepatitis C virus (HCV) RNA sequence at the base position given in the descriptor line. The HCV sequence was screened for optimal ribozyme target sites using a computer folding algorithm and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised to target these sites and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes of the invention inhibit gene expression and/or viral replication, and are used to treat diseases associated with Hepatitis C virus (HCV) infection, e.g. cirrhosis, liver failure and hepatocellular carcinoma. The ribozymes may be used in combination with interference to the infection, e.g. cirrhosis, liver failure and interference to the infection of the combination with the patities of the combination with the combination with the combination of the combination with the com
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18-SEP-1998;
25-FEB-1999;
                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                     hepatocellular carcinoma. The ribozymes may be used in combin interferon to treat HCV infection, other infectious diseases, diseases, and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel ribozymes for the treatment hepatitis C infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-NOV-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Enzymatic nucleic acid; hammerhead ribozyme; virus replication; cleavag cirrhosis; liver failure; hepatocellular carcinoma; interferon; cancer; autoimmune disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ64816 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                W09955847-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-MAR-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-)
                                                                                                             1495
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1488 GAAGCCAGACTTC 1500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RIBOZYME PHARM
                                                                                                        GACTTCAGCAGCCA 1508
                                                                                                                                                                                                                                                                    14 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 98; 123pp; English.
                                                            GAGTTGAGCAGCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        for hairpin ribozyme which cleaves HCV at nt.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mcswiggen
                                                                                                                                                              Conservative
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98US-0100842P.
99US-00257608.
99US-00274553.
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                                                                                                                                                                                                                                                                 2 A; 5 C; 3 G; 0 T; 4 U; 0 Other;
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85.7%;
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                                                                                                                                                                                       Score 10.8;
Pred. No. 27;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                             Length 14;
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                                                                                                                                                                                                                                                                                                                                                autoimmune
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ABX01653/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                     Enzymatic nucleic acid; RNA cleavage; Hepatitis C virus infection; HCV ribozyme; HCV expression; HCV replication; cirrhosis; virucide; liver failure; hepatocellular carcinoma; HCV infection; drug therapy; type I interferon; interferon alpha; interferon beta; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to methods for producing plants resistant to geminivirus, involving introducing a geminivirus replication associated protein (Rep)-iteron antagonist into a plant. The antagonist is a nucleotide sequence of a geminivirus iteron capable of binding to a Rep protein or a defective Rep which has a conserved geminivirus iteron binding site. The present sequence is a geminivirus Rep repeat motif sequence. (Updated on 15-SEP-2003 to standardise OS field)
                                              type I interferon;
interferon gamma; c
substrate; hairpin
                                                                                                                                                                                                                                                                                                                                              ABX01653;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 14 BP; 1 A; 3 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Producing plants resistant to geminivirus, and inhibiting geminivirus replication in plants, by introducing replication associated protein iteron antagonists into the plant, plant cells or propagules.
  Hepatitis C virus.
                                                                                                                                                                                                                                      Hepatitis C
                                                                                                                                                                                                                                                                                            23-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                  ABX01653
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25-JAN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-JAN-2000; 2000WO-US001849
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SCRI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1487 AGAAGCCAGACTTC 1500
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12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGACGCCAGACTCC
                                                                                                                                                                                                                                      virus substrate #138 for HCV hairpin ribozyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chatterji A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rep repeat motif sequence
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                                                                            consensus interferon; hepatotropic; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                               RNA;
                                                    ribozyme; HP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           36.0%;
85.7%;
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Pred. No. 27
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Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ribozymes are useful for modulating the expression and/or replication of HCV. They can be used to treat cirrhosis, liver failure and/or hepatocellular carcinoma. The HCV ribozymes are also useful for treating a condition associated with HCV ribozymes are also useful for core other drug therapies, particularly type I interferon, especially interferon alpha, beta or gamma or consensus interferon. The present sequence represents a substrate for a HCV hairpin (HP) ribozyme. Note: Some of the sequence data for this patent did not form part of the printed specification. The complete sequence data for this patent was obtained in electronic format directly from the USPTO web site at
                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to enzymatic nucleic acids which specifically cleave RNA derived from Hepatitis C virus (HCV). The enzymatic nucleic acid or ribozyme is in a hammerhead (HH) or hairpin (HP) motif where the binding arms comprise sequences complementary to one of the substrate sequences defined in the specification. The HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New ribozymes targeting RNA derived from hepatitis C virus inhibit viral replication and are useful to treat hepatitis C virus infections and cirrhosis, liver failure or hepatocellular carcinoma.
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                                                                                                                                                                                                                                                                                               Oligonucleotide SEQ
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(PAVC/)
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                            18-OCT-2001
                                                                                 WO200177384-A2
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                                                                                                                                                                                                                                                                                                                                                                                                           ABF18234;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABF18234 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      seqdata.uspto.gov/psipsDIDEntry.html
                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Page 62; 80pp; English.
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PAVCO P A.
MACEJACK D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BLATT L.
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85.7%;
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RESULT 24
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XX ABF118
XX SNP;
XW SNP;
KW Centl
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PT Glesi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                               Olek A,
                                                                                                                                                                                                 06-APR-2001; 2001WO-IB000713
                                                                                                                                                                                                                                                                                                                                 central nervous
                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human;
peptide nucleic acid; cytosine methylation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
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                                                                WPI; 2001-657177/75
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                                                                                               Piepenbrock
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Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin

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RESULT 25
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19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                            cytotoxic T-lymphocyte; tumour antigen; immunotherapy; anticancer; ss
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19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                     SAGE tag; serial analysis of gene expression; antigen-presenting ceAPC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL;
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98US-0090040P.
98US-0090041P.
98US-0090042P.
98US-0090043P.
98US-0090044P.
98US-0090047P.
98US-0090047P.
98US-0090047P.
                                                                                                               98US-0090000P.
98US-0090035P.
98US-0090036P.
98US-0090039P.
                                                                                                                                                                   SU86
                                                                                                                                                                              98US-
                                                                                                                                                                                             SN86
                                                                                                                                                                                                      98US-0089993P.
                                                                                                                                                                                                                                98US-0089878P
98US-0089991P
                                                                                                                                                                                                                                                                   98US-0089833P
98US-0089844P
                                                                                                                                                                                                                                                                                                           99WO-US013800
                                                                                                                                                                              -0089997P
                                                                                                                                                                   96666800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 34.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 10.4;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     oligonucleotide primers or peptide nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO:1327.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                 cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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RESULT 26
AAZ83653/c
ID AAZ836
XX
AC AAZ836
XX
AC AAZ836
XX
DT 07-APR
XX
DE Metast

AAZ83653 standard; DNA; 10

07-APR-2000 AAZ83653;

(first entry)

Metastatic breast tumour cell upregulated transcript tag #2887.

멍 S

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AGAAGCCAGA

10

Query Match
Best Local Similarity
Matches 10; Conserv

Conservative

33...; 100.0%; Pr

Mismatches

0

Gaps

0

33.3%; Score 10; 100.0%; Pred. No.

DB 1; Length 10; 28;

1487 AGAAGCCAGA 1496

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complex) and subsequent recognition by T-cell receptors is alone insufficient to activate a robust cytotoxic immune response that can lyse the tumour cells, immunostimulatory cofactors also being required for efficient activation of cytotoxic T-lymphocytes (CTLs). Nucleic acid sequences identified using the SAGE tags have several potential uses. They may be used in vaccines to induce an immune response, particularly against a tumour antigen; to modulate the genotype of an APC; to screen for agents that modulate expression of differentially expressed genes in an APC; and as hybridisation probes/amplification primers for the diagnosis, prognosis and monitoring of diseases related to abnormal expressed genes, or of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing these genes can be used in active immunotherapy (or to stimulate production of a population of antigen-specific effector cells) and vectors containing these genes and APC-associated costimulatory factors ensures adequate antigen and APC-associated costimulatory factors ensures after the APC for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         with monocytes. Some of the transcripts correspond to known genes or ESTs (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while other transcripts correspond to novel genes. Antigen-presenting cell (APC)-associated costimulatory factors play an important role in the activation of the cytotoxic immune response, particularly against tumour cells. Tumour antigen presentation via the MHC (major histocompatibility
Sequence
                                          presentation to endogenous APCs and upregulates the APCs for the presentation of co-stimulatory signals, migration to T cell-rich secretion of T cell growth factors and secretion of chemokines for recruitment of immune effector cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequences AAZ77573-Z79709 represent SAGE (serial analysis of gene expression) tags used to identify mRNA transcripts encoding immunostimulatory cofactor proteins which are preferentially or differentially expressed in monocyte-derived dendritic cells compared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cells, useful in gene vaccines against cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated polynucleotides differentially expressed in antigen-presenting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-106077/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SHAN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GENZ)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ROBE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENZYME CORP.
ROBERTS B L.
SHANKARA S.
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B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shankara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98US-0090076P.
98US-0090077P.
98US-0090078P.
98US-0090079P.
98US-0090080P.
98US-0111715P.
v
Α;
2 C; 3
G; 0 T; 0
  u;
o
  Other;
                                                                for
                                                                                    sites,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or ESTs
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AAF39527
ID AAF3
XX
AC AAF3
XX
DT 23-1
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                                                                                                                                                                                                                                                                                                                  AZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour CC tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 CC to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour cells). These CC transcripts can be used for diagnosis, prognosis, monitoring and CC treatment of breast cancer, particularly where metastatic Diagnosis is by standard immunoassays or hybridisation/amplification reactions. CC Compounds that modulate expression of the transcripts are potentially CC useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of c.g. therapeutic genes (also ribozymes or antisense sequences), CC particularly an antigen-encoding sequence for use in gene or cell-based concines. Polypeptides encoded by the transcripts are also useful in CC vaccines; for diagnosing breast cancer and for raising specific capents. Host cells that produce the polypeptides or as therapeutic and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic I lymphocytes, and these used for adoptive
                                                                                               RESULT 27
                                                                                                                                                                                                                              Query Match
Best Local :
                                                                                                                                                                                                             Best Local Similarity Matches 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Isolated polynucleotides differentially expressed between metastatic non-metastatic breast cancer cells, useful for diagnosis, prevention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9965928-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                 AAF39527;
                                                                                                                                                                                                                                                                           Sequence 10 BP; 2 A; 2 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 136; 219pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treatment of cancer.
 23-MAR-2001
                                                                 AAF39527 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-106079/09
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENZ )
                                                                                                                                                                               1496 ACTTCAGCAG 1505
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ROBERTS B L.
SHANKARA S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BL,
                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shankara
 (first entry)
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98US-0089997P.
98US-0090039P.
98US-0090040P.
98US-0090041P.
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                                                                                                                                                                                                                          DB 1; Length 10; 28;
                                                                                                                                                                                                               0
                                                                                                                                                                                                               Indels
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Yeast NORF gene SACE tag oligonucleotide SEQ ID NO:6266.

Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
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Saccharomyces cerevisiae.

linker; PCR primer; ds.

WO200077214-A2.

21-DEC-2000.

14-JUN-2000; 2000WO-US016223.

16-JUN-1999; 99US-00335032.

(UYJO) UNIV JOHNS HOPKINS.

Velculescu V, Vogelstein B, Kinzler K;

WPI; 2001-061874/07

Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.

Example; Page 223; 419pp; English.

Coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not comprising a ssigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also codescribed are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at cleast 10% between any two phases of the cell cycle selected from log cantifungal drugs comprising: (a) a method (M2) for screening candidate cell; and (b) monitoring expression of a NORF gene whose expression of varies as in M1, where a test substance which modifies the expression cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human DNA with a probe which comprises at least 10 comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; cell cycle progression in the yeast cell of at least 1 NORF gene whose expression in a class of drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell with a candidate drug and member of a cycle and (d) a method (M4) for identifying a candidate drug as a member of a cycle and genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle. The correspondence of the cell cycle with a candidate of the cell cycle. The correspondence of the cell cycle is the cell cycle. The cycle and for identification of the present invention.

Apr33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.

Sequence 10 BP; 5 A; 2 C; 3 G; 0 T; 0 U; 0 Other;

밁 Ś Matches Query Match Best Local Similarity 1487 10; AGAAGCCAGA 1496 Conservative 100.0%; 33.3%; 0; Score 10; Pred. No. Mismatches DB 1; Length 10; 28; 0; 0

Gaps

0

RESULT 28
AAF34162
ID AAF34162 st.
XX

AAF34162 standard; DNA; 10 BP.

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                                    δ
                                                                                                                                                                                                           The present invention describes an isolated DNA molecule comprising a CC coding sequence of a yeast gene selected from a group of 745 NORF (not CC previously assigned open reading frame; or nonamnotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC comprising a SAGE (serial analysis of gene expression) tag. Also CC cleast 10% between any two phases of the cell cycle selected from log CC phase, S phase and G2/M; (2) a method (M2) for screening candidate CC ell; and (b) monitoring expression of a NORF gene whose expression of CC varies as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for CC identifying human genes which are involved in cell cycle progression of CC comprising contacting human DNA with a probe which comprises at least 10 CC contiguous nucleotides of a NORF gene whose expression of CC class of drugs having a characteristic effect on gene expression in a CC class of drugs having a characteristic effect on gene expression in a CC expression in the yeast cell of at least 1 NORF gene whose comprises and affect phases of drugs. The NORF gene whose capression in a GC expression in the yeast cell of at least 1 NORF gene whose capression is affected by the class of drugs. The NORF gene whose capression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially cycle and for identification of antifungal drugs which affect the cell cycle. The CC expressed genes may be used as markers of phases of the present invention. CC AAR33267 to AAR33267 represent in the exemplification of the present invention. CC AAR33261 to AAR33267 represent invention.
                                                                                 Matches
                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                                                                            Sequence 10 BP; 5 A; 2 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-JUN-2000; 2000WO-US016223.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-MAR-2001 (first entry)
                                      1486 AAGAAGCCAG 1495
                                                                               10;
ب
                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 32; 419pp; English.
                                                                                                                                                                                                   the exemplification of the present invention
                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9908-00335032
                                                                                                100.0%;
                                                                                                                     33.3%;
                                                                           Score 10; DB;
; Pred. No. 28;
0; Mismatches
                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kinzler
                                                                                                    DB 1;
. 28;
                                                                             0
                                                                                                                 Length 10
                                                                             0
                                                                           Gaps
                                                                             0
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RESULT 29

밁 Ş

Matches

10;

Conservative

Similarity

100.0%; 33.3%;

Score 10; Pred. No.

DB 1; Length 10, 28;

Mismatches

0;

Indels

0

Gaps

0

1496 ACTTCAGCAG 1505

Sequence 10

BP; 3

A; 3 C; 2

G; 2 T; 0 U; 0 Other;

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The present invention describes an isolated DNA molecule comprising a CC coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also Cd described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log thase, S phase and G2/M; (2) a method (M2) for screening candidate CC antifungal drugs comprising; (a) contacting a test substance with a yeast CC varies as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human genes which are involved in cell cycle progression of comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a cyeast cell of at least 1 NORF gene whose expression in a cyeast cell of at least 1 NORF gene may be used to study, monitor and affect phases of the cell cycle, the differentially cycle and for identify candidate drugs. The NORF gene may be used to study, monitor and affect phases of the cell cycle. The ceptes may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF33268 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-061874/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Velculescu V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yeast NORF gene SAGE tag oligonucleotide SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example; Page 221; 419pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-JUN-2000; 2000WO-US016223
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-00335032
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kinzler K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IJ
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RESULT 30
AAF41899
                                                                                       Matches
                           Best Local Similarity
                                                                                                                     comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame, or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-061874/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Velculescu V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200077214-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:8638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example; Page 308; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-JUN-2000; 2000WO-US016223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-DEC-2000
                                                                                     Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-JUN-1999;
        10;
                                                                                       BP;
      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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33.5°.
100.0%; F1
                                                                                     2 C; 3 G; 0 T; 0 U; 0 Other;
                         33.3%; Score 10; 100.0%; Pred. No.
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        Mismatches
                           DB 1; Length 10; 28;
        0
        Indels
        0
        Gaps
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RESULT 32
AAX89228
ID AAX89
XX
AC AAX89
XX
DT 15-SE
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                                                                                                                                                                                          Best Local Similarity Matches 10; Conserv
                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                       expression by random primed reverse transcription PCR (RT-PCR). The primer sequences are generated by stimulating PCR reactions on non-redundant mammalian nucleotide sequence databank entries containing at least 1,000 bp of coding region. The primers selected, such as the present one, had to meet various criteria such as having an efficiency index between 2-10, having a selectivity index higher than 1, being 12 bp long i.e. 8 C or G and 4 T or A, and each primer differed from the others in at least 5 of the 8 bases at the 3'-end. The invention claims the selected primers make it possible to use internally primed, PCR-based RNA fingerprinting for simple, exhaustive and systematic analysis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Random primed reverse transcription PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV32260
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV32260 standard; DNA; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Differential screening of gene expression by reverse transcription polymerase chain reaction - uses random priming with primers selected for high efficiency and selectivity by computer screening of database(s).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-APR-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                differential gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RT-PCR; primer; amplification; reverse transcription; RNA fingerprinting;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-AUG-1998
                           AAX89228;
                                                                                                                                                                                                                                                    Sequence 12 BP; 0 A; 3 C; 4 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                              coding
                                                                                                                                                                                                                                                                                           differential gene expression as an advantageous alternative to differential display. The method can also be useful for isolat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention provides a method for the differential screening of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 9; Page 24; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-230725/20..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Consalez G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-SEP-1997;
                                                         AAX89228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SANR-) FOND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1482
                                                                                                                                                           1479 CACGACCAAG 1488
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                                                                                                                                                                                                                                                                               sequences and to compare
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GACCAAGAAG
                                                         standard; DNA; 13
                                                                                                                                 CACGACCAAG
                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fesce R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CENT SAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96GB-00020216.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97WO-EP005290
                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.0
                                                                                                                                                                                                                       33.3%;
                                                                                                                                                                                                                                                                                 The method can also be useful to compare known and new genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RAFFAELE DEL MONTE TABOR
                                                           ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВÞ
                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                         Score 10;
Pred. No.
                                                                                                                                                                                                          DB 1;
33;
                                                                                                                                                                                                                                                                               new genes
                                                                                                                                                                                             0
                                                                                                                                                                                                                       Length 12
                                                                                                                                                                                             Indels
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15-SEP-1999

(first entry)

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                                                                                                                                                                                                                                                                                                                                                                                                    The invention provides an antibody specific for epitopes (AAY27164-175) located on the extracellular domain of STZL protein. The antibody is used: (a) in a binding assay to identify and/or distinguish T-helper 1 (Th1) or Th2 and evaluate the abundance of Th2 in a sample of body fluid or tissue; (b) to lyse Th2 cells in preference to Th1 cells or otherwise inhibit Th2 cell function; and (c) as a medicament for use in therapy as a mediator of an allergic reaction, e.g. asthma; diseases mediated by intracellular pathogens, especially arthritis. The antibody is targeted to an epitope (STZL) which is a member of the immunoglobulin superfamily and is found on Th2 cells and not Th1 cells
                                                                                              Synthetic
                                                                                                                      Camellia sinensis L. (O.) Kuntze;
                                                                                                                                              Camellia
                                                                                                                                                                                                                     ADC64953 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                Sequence 13 BP; 4 A; 4 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
          30-MAR-2001; 2001CN-00112459
                                   06-NOV-2002
                                                           CN1377966-A
                                                                                  Camellia sinensis.
                                                                                                                                                                      18-DEC-2003
                                                                                                                                                                                            ADC64953;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 12; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-430265/36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Liew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ST2L; extracellular domain; mouse; epitope; antibody; T-helper cell; Th1 Th2; medicament; therapy; allergic reaction; asthma; Leishmania; incracellular pathogen; inflammatory disease; arthritis; immunoglobulin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer H-AP34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UNIU)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ovel antibody to ST2L, helper 1 cells.
                                                                                                                                                                                                                                                                                                         1497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     primer; ss.
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                                                                                                                                                                                                                                                                                               CTTCAGCAGC 1506
                                                                                                                                              sinensis L.
                                                                                                                                                                                                                                                                                 CTTCAGCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Xu D;
                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                     (first
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                                                                                                                                                                     entry)
                                                                                                                                              (O.) Kuntze
                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                         33.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           an epitope found on
                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                            Score 10; DB Pred. No. 36;
                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                   tea tree; PCR primer; ss
                                                                                                                                          related PCR primer AP34.
                                                                                                                                                                                                                                                                                                                                                        DB 1;
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                                                        Query Match
Best Local Similarity
Matches 10; Conserv
                                                                                                                  Sequence 13
                                                                                                                                                                         The present invention describes the cloning of a new gene sequence expressed and inhibited during hibernation of the top plumelet of camellia sinensis L.(O.) Kuntze tea tree. Also described is the identification, cloning and analysis of a primer terminal in the grequence expressed and inhibited during hibernation of the top pluments.
                                                                                                                                                                                                                                                                                          Cloning of a new gene sequence expressed and inhibited during winter dormancy of a tea tree top plumelet, comprises identification, cloning and analysis of a new primer in the gene sequence.
                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                                                                                                                                                                                Example 3; Page 32; 66pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                30-MAR-2001; 2001CN-00112459.
                                                                                                                                    the tea tree. The present sequence represents a PCR primer which is a in an example from the present invention.
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                CTTCAGCAGC 1506
 CTTCAGCAGC
                                                           Conservative
                                                                                                                  BP; 4 A; 4 C; 3 G; 2 T; 0 U; 0 Other;
13
                                                                                     33.3%;
                                                                         100.0%;
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                                                                       Score 10;
Pred. No.
                                                           Mismatches
                                                                       36;
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                                                                                     Length 13
                                                           Indels
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                                                                                                                                                                                          the gene
                                                         Gaps
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RESULT 34
ABH65692/c
ID ABH656
XX ABH656
XX ABH656
XX SNP; E
KW Peptid
KW Centra
XX Homo E
XX HO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           single nucleotide polymorphism; human; diagnosis; PNA; cancer;
ide nucleic acid; cytosine methylation; cardiovascular; primer;
ral nervous system; gastrointestinal; respiratory; immune; metal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ID NO 265669 for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВÞ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory,

central nervous system, cardiovascular and

Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and

typing, i cytosine

methylation

WPI; 2001-657177/75.

07-APR-2000;

2000DE-01019173

(EPIG-)

EPIGENOMICS

ĀG

Piepenbrock C,

Berlin

06-APR-2001; 2001WO-IB000713.

Claim 1; SEQ ID NO 265669; 29pp + Sequence Listing; German.

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RESULT 35
ABH65694/c
ID ABH656
XX ABH666
XX AB
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Best Local S
Matches 11
   Matches
                                       Best Local
                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                        acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABF00010-ABF9989,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                 represent the oligomers described in the invention. NOTE: The sequent data for this patent did not form part of the printed specification, was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13 BP; 3 A; 0 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 265671; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Olek A,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 13 BP; 6 A; 5 C; 0 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
            19-MAR-2002; 2002WO-US008546
                                                                                                                             synthesis;
                                                                                                                                              Molecular function;
                                                                                                                                                                       DNA-templated synthesis related oligonucleotide
                                                                                                                                                                                                        27-JAN-2003
                                                                                                                                                                                                                                    ABQ83680
                                                                                                                                                                                                                                                                 ABQ83680 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 265672; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-657177/75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-APR-2000; 2000DE-01019173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-APR-2001; 2001WO-IB000713.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200177384-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide SEQ ID
                                         26-SEP-2002.
                                                                     WO200274929-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (EPIG-)
                                                                                                                                                                                                                                                                                                                                                                    1489 AAGCCAGACTTCA 1501
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                                                                                                                                                                                                                                                                                                                                          AACCCAAACTTCA 13
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                                                                                                                             library;
                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                      (first entry)
                                                                                                                            diversification; selection; amplification; evolve; chemical compound; hybridisation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                 32.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              265672 for detecting SNP TSC0064388.
                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                Score 9.8; D
Pred. No. 39;
                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 13;
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                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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RESULT 39
ABV70346/c
ID ABV703:
XX ABV703:
XX ABV703:
XX 21-OCT
XX 21-OCT
XX Human;
XX Human;
XX Human;
XX Human;
XX Homo 8
XX 980718
XX WO2002
XX WO2002
XX PD 11-JUL
XX PF 20-DEC
XX

Human skin EST 8132.

21-OCT-2002

(first

ABV70346;

ABV70346 standard; cDNA; 11 BP

immunosuppressive; antiinflammatory; cytostatic; SAGE; neuro psoriasis; dermatitis; skin cancer; EST; expressed sequence

Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis;

20-DEC-2001; 2001WO-EP015179

11-JUL-2002.

sapiens.

밁 S

13

GGCAGATTTCAGC 1

Query Match
Best Local Similarity
Matches 11; Conserv

Conservative

0

32.7**%**; 84.6**%**;

Score 9.8; I Pred. No. 39; Mismatches

ĎΒ

1; Length 13 Indels

٥,

Gaps

0

1491 GCCAGACTTCAGC 1503

Sequence 13 BP; 3 A; 4 C; 3 G; 3 T; 0 U; 0 Other;

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contacting one or more transfer units having an anti-codon and reactive unit with the one or more templates under conditions to allow for hybridisation of the one or more anti-codons to template, and reaction of the reactive units. Also described: (1) a method (M2) of evolving a library of compounds; (2) a kit comprising one or more nucleic acid templates and one or more non-natural polymers; and (4) a library (1) comprising one or more non-natural polymers; and (4) a library (1) compounds is bonded to an amplifiable template whose nucleotide sequence is informative of the structure of the chemical compounds where the library is synthesised by M1 or M3. The method can be used for synthesising one or more chemical compounds which contain an anti-codon comprising a nucleotide sequence which hybridises with one or more nucleic acid templates. M1 is useful for synthesising a library of chemical compounds that are not, or do not resemble nucleic acids or nucleic acid analogues. The present sequence represents an oligonucleotide which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthesizing chemical compounds by hybridizing one or more templates which have associated reactive unit, with one or more transfer units having anti-codon and reactive unit, and performing reaction of reactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-MAR-2001; 2001US-0277081P.
19-MAR-2001; 2001US-0277094P.
20-JUL-2001; 2001US-0306691P.
19-MAR-2002; 2002US-00101030.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Fig 32; 146pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Liu
                                                                                                                                                                                                                                                                                                                                                                                                                                                  more chemical compounds. M1 involves providing one or more temp. which optionally have a reactive unit associated with them; and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        units.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present invention describes a method (M1) for synthesising one or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gartner ZJ,
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03-JAN-2001; 2001DE-01000127.

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RESULT 40
ABV69357/C
ID ABV69357/C
ID ABV69357/C
ABV69357/C
XX
ABV69357/C
XX
ID Human
XX
Immuni
XW
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Immuni
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Immu
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SACE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 11 BP; 1 A; 3 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-590638/63
                                                        e.g. skin cancer
                                                                                   In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                             WPI; 2002-590638/63
                                                                                                                                                                                                                                                                                                                                                    03-JAN-2001; 2001DE-01000127
                                                                                                                                                                                                                                                                                                                                                                                                          20-DEC-2001; 2001WO-EP015179
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human skin EST 7143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABV69357 standard; cDNA; 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                     Conradt
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in the skin of humans or animals by subjecting a mixture of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
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Sequence 11 BP; 0 A; 4 C; 2 G; 5 T; 0 U; 0 Other;

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밁
              Ş
                                Query Match
Best Local S
Matches 10
                 1485 CAAGAAGCCAG 1495
11 CAAGAAGGCAG 1
                                  10;
                                         Similarity
                                  Conservative
                                         31.3%;
                                  0
                                         Score 9.4;
Pred. No. 4
                                  Mismatches
                                                   DB 1;
                                  1,
                                                 Length 11;
                                  Indels
                                  <u>.</u>.
                                  Gaps
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0

ARESULT 41
ABV6295/C
ID ABV629
XX ABV629
XX ABV629
XX DT 21-OCT
XX Human;
XX Human;
XX Human;
XX Homo s
XX WO2002
XX II-JUL
XX O3-JAN
XX O3-JAN
XX PA (HENK
XX PA (HENK
XX PA (HENK
XX PE ers
PT e.g. s
DR WPI; 2
XX II vit
PT homeos
PT e.g. s
CC in the
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CC encode
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CC skin. Human skin EST 21-OCT-2002 ABV62925; ABV62925 standard; cDNA; 11 (first entry) 711. ВP

immunosuppressive; antiinflammatory; cytostatic; SAGE; neuro psoriasis; dermatitis; skin cancer; EST; expressed sequence Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; tag; ss.

Homo sapiens.

WO200253774-A2

11-JUL-2002

20-DEC-2001; 2001WO-EP015179

03-JAN-2001; 2001DE-01000127

'n Conradt M, Hofmann <u>~</u>

2002-590638/63

homeostasis and identifying In vitro identification of skin-expressed .g. skin cancer. cosmetic မ္ပ ed genes, useful pharmaceutical a agents for determining

Disclosure; Page 45; 1345pp; German.

The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression (M1) is useful for identifying genes and quantify their expression (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis, atopic dermatitis; acne; seborrhea; lupus erythematosus; rosaces; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention

Disclosure; Page 224; 1345pp; German

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RESULT 42
ABV68643;
ID ABV68643;
XX Human; ski
XX WO20025377
XX WO20025377
XX WO20025377
XX WO20025377
XX WO20025377
XX WFI; 2002-
XX WFI;
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Best Local S
Matches 10
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Best Local Similarity
Matches 10; Conserv
                                                                                                                                                                                                                                                                                                           The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis are to to test agent (A) that maintains or promoters, specifically neurodermatitis; sunburn, psoriasis; scleroderma; ichthyosis; atopic dermatitis; ane; sebornhea; lupus erythematosus; robacca; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
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                                                                                                                                                                                                                               Sequence 11 BP; 3 A; 2 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 204; 1345pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-JAN-2001; 2001DE-01000127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human skin EST 6429
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                                                       1489 AAGCCAGACTT 1499
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12
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AAGCCAGTCTT 1
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                                                                                                                 Conservative
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                                                                                                                                            31.3%;
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                                                                                                                 0
                                                                                                                                                                                                                               G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 9.4;
Pred. No.
                                                                                                                                            Score 9.4;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                 Mismatches
                                                                                                                                                                     9.4;
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                                                                                                                                                                     DB 1; Length 11;
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                                                                                                              1;
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                                                                                                              Gaps
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RESULT 43

polymer;

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ABV69164
XX ABV6
XX ABV6
XX ABV6
XX ABV6
XX Huma
XX Huma
XX Huma
XX Homc
XX Hom
RESULT 44
AAZ41704/c
ID AAZ4177
XX AAZ417
XX AAZ417
XX DT 20-MAR
DT 21-JAN
XX Organi
XX Amplif
KW detect
KW organo
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Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; sebornhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
Amplification; polymerase chain reaction; PCR; microorganism; detection; pollutant; soil; food; agricultural chemical; polynorganochlorine; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABV69164 standard; cDNA; 11 BP
                                                                                                                       Organic material detecting primer 65.
                                                                                                                                                                                    20-MAR-2003
21-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 11 BP; 7 A; 1 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 218; 1345pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               e.g. skin cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-590638/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-JAN-2001; 2001DE-01000127.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human skin EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABV69164;
                                                                                                                                                                                                                                                                                 AAZ41704;
                                                                                                                                                                                                                                                                                                                                        AAZ41704 standard; DNA; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-DEC-2001; 2001WO-EP015179.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HENK ) HENKEL KGAA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1486 AAGAAGCCAGA 1496
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H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAGAAGCAAGA 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the invention
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                                                                                                                                                                                    (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31.3%;
90.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hofmann
                                                                                                                                                                                                                                                                                                                                               BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 9.4;
Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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RESULT 45
AAZ41488/c
ID AAZ414
XX AAZ414
XX Hicrob
XX Microb
XX Microb
XX Microb
XX Microb
XX Fandom
XX Synthe
XX Synthe
XX JP1127
XX JP1127
XX JP1127
XX J1-MAR
XX SAOL
PA (SAOL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 comprising (i) preparing many primers (P) with different probabilities of amplification and (ii) simultaneous polymerase chain reaction (PCR) of many different DNA using these primers. The method is used (i) to differentiate between different microorganisms in a mixed population and (ii) to determine presence/absence of an impurity (pollutant), or its concentration, in e.g. soil, foods, compost etc., typically metals, agricultural chemicals, polymers, organochlorine compounds etc. A particular use is monitoring composting of organic material.

Amplification with many primers produces a lot of information, so reliability of the test is improved, and many samples may be tested
                                                                                                                                                                         Microbe; detection; organic waste; arbitrarily primer PCR; random amplified polymorphic DNA; amplification; PCR prime
                                                                                                                    JP11276176-A
                                                                                                                                               Synthetic
                                                                                                                                                                                                                      Microbe
                                                                                                                                                                                                                                                 19-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 12 BP; 1 A; 3 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        quickly. AAZ41640-Z41855 represent the primers described the invention. (Updated on 20-MAR-2003 to correct PR fiel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      microorganisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SAOL )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-OCT-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                               31-MAR-1998;
                                                           31-MAR-1998;
                                                                                       12-OCT-1999.
                                                                                                                                                                                                                                                                                                          AAZ41488 standard; DNA; 12.BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes a novel method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-592157/51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-MAR-1999;
    (SAOL )
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                           1484 CCAAGAAGCCA 1494
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polymerase chain reaction method, for differentiating between organisms and for detecting contaminants.
                                                                                                                                                                                                                                                                                                                                                                                                                                      10;
                                                                                                                                                                                                                                                                                                                                                                              11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SANYO ELECTRIC CO LTD. SOC TECHNO-INNOVATION
    SANYO
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                     detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TECHNO-INNOVATION AGRIC FORESTY & FI
                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                (first entry)
    ELECTRIC CO LTD
                               98JP-00087652.
                                                             98JP-00087652
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98JP-00087651.
99JP-00069694.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99DE-01014461
                                                                                                                                                                                                                     in organic waste arbitrarily primed PCR primer #65
                                                                                                                                                                                                                                                                                                                                                                                                                                                 31.3%;
90.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 9.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 43;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for the amplification of DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PR field.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                         primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of.
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A method has been developed for the amplification of a DNA fragment in which amplification is carried out on the DNA fragments of a number of different DNAs. The method comprises a PCR reaction repeatedly carrying out a heat-denaturing step, a primer annealing step and a polymerase extending step, to amplify the DNA fragments of a plural of different DNAs. The method can detect the existence of a microbe in organic waste. AAZ41424 to AAZ41639 represent PCR primers used in random amplified polymorphic DNA arbitrarily primed PCR, for the detection of microbes in
                                                                                                                                                                                                                                 Example; Page 9; 40pp; Japanese
                                                                                                                                                                                                                                                                        Amplification of a DNA existence of a microbe.
                                    organic waste
                                                                                                                                                                                                                                                                                                                                                                         (NORI-) ZH NORIN SUISAN
                                                                                                                                                                                                                                                                                           fragment
                                                                                                                                                                                                                                                                                                                                                                         SENTAN GIJUTSU SANGYO
                                                                                                                                                                                                                                                                                               'n
                                                                                                                                                                                                                                                                                             order to establish the
                                                                                                                                                                                                                                                                                               of.
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밁 á Matches Query Match Sequence 12 BP; 1 A; 3 C; 4 Local 1484 CCAAGAAGCCA 1494 l Similarity 10; Conserv CCAAGAGGCCA 1 Conservative 31.3%; G; 4 T; 0 U; 0 Other; Score 9.4; Pred. No. 4 Mismatches DB 1; Length 12; Indels ٥,

0

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RESULT 46
AAC97839/c
Synthetic
                                                                                                                                            28-FEB-2001
                                                                                                                                                    AAC97839;
                                                                                                                                                            AAC97839
                                                                                                     03-OCT-2000
                                                                                                             JP2000270867-A
                                                                                                                            Primer; amplification; selective; ss.
                                                                                                                                    Primer used
                                                                                                                                                            standard;
                                                                                                                                    to illustrate DNA amplification method SEQ ID
                                                                                                                                            (first entry)
                                                                                                                                                            DNA;
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WPI; 2001-011047/02 19-MAR-1999; 19-MAR-1999; (SAOL) (NORI-) SANYO ELECTRIC CO LTD. ZH NORIN SUISAN SENTAN 99JP-00076844. 99JP-00076844. GIJUTSU SANGYO

of a DNA fragment and ite apparatus

Example 1; Page 8; 22pp; Japanese.

method comprises successive repetitions of heat-denaturing, annealing of a primer and an extending step using a DNA polymerase. The method makes use of a cDNA pool in which the primer is one primer or a pair of primer sets and has an amplification probability which allows it to amplify a DNA fragment from a limited number of the cDNAs among the DNA pool (where the limited number is in the range of 1 to 25). Also included in the invention are apparatus used for carrying out the method, a primer and a DNA polymerase and a kit used for amplifying a DNA fragment. The method can be used to amplify a limited number of cDNAs from a pool in which a wide variety of cDNAs are present. Oligonucleotides AAC97775 - AAC97990 represent primers used in an example illustrating the method of the invention relates to a method for amplifying a DNA fragment. The

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RESULT 47
ABIO2085/c
ID ABIO20
XX ABIO20
XX Oligon
XX SNP; B
KW Peptid
KW Centra
XX SNP Homo B
XX Homo B
XX Homo B
XX Homo B
XX WO2001
XX D6-APR
XX O1-APR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
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                                                                                                                  Oligonucleotide primer SEQ ID NO 353642 for detecting SNP TSC0048625.
                                                                                                                                                                                22-FEB-2002
                                                                                                                                                                                                                                       ABI53669;
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                                                                                                                                                                                                                                                                                                  ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>.</u>.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 43;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 9.4;
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metab

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RESULT 50
AA199727/c
ID AA1997
XX AA1997
XX AA1997
XX AI997
XX Microb
XX Microb
XX PCR pr
KW Arseni
XX Synthe
PN W02001
XX W02001
XX 11-OCT
XX 11-OCT
XX 27-MAR
XX 15-MAR
PR 31-MAF
PR 31-MAF
PR (SAOL
PA (NORQ
XX
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Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cyrosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          designed to detect methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-APR-2000; 2000DE-01019173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-APR-2001; 2001WO-IB000713.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200177384-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo
                                                                                                                                                                                                        PCR primer; microbe; composted waste; soil; contaminant; mercury; arsenic; dioxin; hormone; SSC-PCR;.ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 12 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 353642; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Set of oligonucleotides, useful for diagnosis and cell typing, is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-OCT-2001.
                                                          31-MAR-2000; 2000JP-00099482.
                                                                                       27-MAR-2001; 2001WO-JP002516
                                                                                                                                                  WO200175156-A1
                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                    Microbial SSC-PCR primer SEQ ID NO 10
                                                                                                                                                                                                                                                                                   21-JAN-2002
                                                                                                                                                                                                                                                                                                                                            AAI99727 standard; DNA; 12 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (EPIG-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                1480 ACGACCAAGAA 1490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention describes novel oligonucleotide primers or peptide nucleic (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                     12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10; Conserv
               SANYO ELECTRIC CO LTD. SOC TECHNO-INNOVATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EPIGENOMICS AG
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                                                                                                                                                                                                                                                                                                                                                                                                                     ACGACCAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 A; 1 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
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                AGRIC FORESTY & FI.
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ARXIB642/c
ID AAXIB6
XX AAXIB6
AC AAXIB6
AC AAXIB6
AC AAXIB6
XX P53; s
XW p53; s
XW rat em
KW tumour
XX Synthe
OS Rattus
XX WO9901
XX WO9901
XX U09901
XX U0
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to identification of microbes, especially for obtaining information on the microbial spectrum in composted wastes and in the soil to which they are to be applied, and especially to give an indication of possible contaminants in the soil (such as mercury, arsenic, dioxins and environmental hormones) by the presence of microorganisms associated with them, comprising: (a) polymerase chain reaction on DNA of a mixture of microbes using a number of primers of different amplification probability (SSC-PCR); (b) electrophoresis of the DNA amplification fragments; (c) detecting the bands obtained on the electrophoresis inage; (d) correcting errors in the image gradients; (e) measuring the position and intensity of the bands obtained on the results and creating a list of the band data; (g) searching the search. The present sequence is that of a PCR primer, useful to the invent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Microbe identification comprises comparing results of reference database for determining microbial spectrum
                              Diagnosis of cancer in potentially neoplastic samples - by comparing the level of transcription between RNA transcripts in two tissue samples, useful for providing an extensive profile of gene expression in rat embryo fibroblast (REF) cells.
                                                                                                                                                                                                                                                                                                                                                                                                                      02-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       p53; serial analysis of gene expression; SAGE tag; cancer; neoplastic; rat embryo fibroblast; REF; tumour suppressor; cell cycle control;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          p53 serial analysis of gene expression tag #45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX18642 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 12 BP; 1 A; 3 C; 4
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                                                                                                                                                                                     WPI; 1999-106079/09.
                                                                                                                                                                                                                                                                                                                                                              02-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9901581-A1
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                                                                                                                                                                                                                                                                                                    (GENZ ) GENZYME CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1484 CCAAGAAGCCA 1494
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                                                                                                                                                                                                                                            Galella EA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                         98WO-US013903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diagnosis; ss.
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                                                                                                                                                                                                                                               Bertelsen AH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВÞ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 9.4;
Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                               Beaudry GA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC A method has been developed for the diagnosis of cancer in potentially CC neoplastic samples. The method comprises comparing the level of CC transcription between RNA transcripts in two tissue samples (which are of CC the same type), where the first sample is potentially neoplastic, and the Second sample is normal human tissue. The first sample is categorized as CC neoplastic if its level of transcription is lower than that of the second CC sample. The transcript is selected from Alu, RAS, UG sanRNA, 16S RNA, EGR-CC 1, ribosomal protein S27, ETS-1, 28S RNA, CGR11, and LIMK-2, and it is cidentified by a tag selected from ribosomal protein L11a, alpha-tubulin CC (T1) and (T2), thymosin beta-4, and gamma-actin. The present sequence represents a serial analysis of gene expression (SAGE) tag from the CC present invention. The use of SAGE tags provides an extensive profile of gene expression in rat embryo fibroblast (REF) cells containing the (non) CC denctional p53 tumour suppression gene. The discovery of new SAGE tags, which are regulated by p53, enables the diagnosis of genes that are contained to cell cycle control and tumourigenesis
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                                        19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                        SAGE tag; serial analysis of gene expression; antigen-presenting cell; APC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL;
                                                                                                                           19-JUN-1998
                                                                                                                                                                                                                                                                                                                            23-DEC-1999
                                                                                                                                                                                                                                                                                                                                                      WO9965924-A2
                                                                                                                                                                                                                                                                                                                                                                                                            cytotoxic T-lymphocyte; tumour antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human dendritic cell SAGE tag,
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                19-JUN-1998;
19-JUN-1998;
                                                                                                                                                       19-JUN-1998
                                                                                                                                                                    19-JUN-1998;
19-JUN-1998;
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                                                                                                                                                                                                                                         19-JUN-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-APR-2000 (first entry)
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S086
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98US-0089844P.
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                                                                                                                                                                                   -0089993P
                                                                                                                                                                                                                            -0089878P
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100.0%; Pred. No. 43;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO:1647
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                                                                                                                                                                                                                                                                                                                                                                                                         immunotherapy; anticancer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 10;
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RESULT 53
AAZ83241/c
ID AAZ832
XX
AC AAZ832
XX

AAZ83241 standard; DNA; 10 BP

AAZ83241;

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10

1500 CAGCAGCCA 1508

Query Match Best Local S Matches 9

Similarity 9; Conserv

Conservative

30.v., 100.0%; F1

30.0%; Score 9; DB: 100.0%; Pred. No. 43 tive 0; Mismatches

Sequence 10

BP; 1

A,

2 C; 4

G; 3 T; 0

U; 0 DB 1;

Other;

Length 10; 0,

Indels

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Gaps

0

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other transcripts correspond to novel genes. Antigen-presenting cells (APC) associated costimulatory factors play an important role in the cativation of the cytotoxic immune response, particularly against tumour cells. Tumour antigen presentation via the MHC (major histocompatibility complex) and subsequent recognition by T-cell receptors is alone insufficient to activate a robust cytotoxic immune response that can lyse the tumour cells, immunostimulatory cofactors also being required for cefficient activation of cytotoxic T-lymphocytes (CTLs). Nucleic acid care gequences identified using the SAGE tags have several potential uses. They may be used in vaccines to induce an immune response, particularly against a tumour antigen; to modulate the genotype of an APC; to screen for agents that modulate expression of differentially expressed genes in an APC; and as hybridisation probes/amplification primers for the capression of these genes. Detection of the dendritic cell differentially expressed genes, or of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing these genes can be used in active immunotherapy (or to stimulate production of a population of antigen specific effector cells) and vectors containing these genes centually expressed genes are used in gene therapy. Co-administration of tumour antigens and corresponded costimulatory factors ensures adequate antigen and presentation of co-stimulatory signals, migration to Tcell-rich sites, secretion of Tcell growth factors and secretion of chemokines for the corresponded proteins.
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
08-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences AA277573-279709 represent SAGE (serial analysis of gene expression) tags used to identify mRNA transcripts encoding immunostimulatory cofactor proteins which are preferentially or differentially expressed in monocyte-derived dendritic cells compared with monocytes. Some of the transcripts correspond to known genes or I (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cells, useful in gene vaccines against cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Roberts BL, Shankara
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(ROBE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and
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                 AAZ83053 standard; DNA; 10 BP
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                                                                                                                             Sequence 10
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                                                                                                  a cell expressing the target epitope with cytotoxic T cells generated against the cell to identify DNA clones expressing the target epitope. The method may also comprise providing a cytotoxic T cell specific for a gene product differentially expressed by a cell and measuring the cross-reactivity of the cytotoxic T cell. The methods are useful for identifying tumour specific target epitopes and antigens which are useful in immunogenic compositions or vaccines to induce the regression of tumours, cancers or infections in mammals. The genes expressed in a panel of tumour cells that are derived from single immortalised, noncells under cell line are used to generate HLA restricted cytotoxic T cells which are evaluated for activity against tumour cells. The method is useful to identify potential antigens expression is altered as a result of infection. The differential gene expression strategies can be applied to identify immunogenic molecules of cells infected with virus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA15239-50 represent arbitrary primers which are used for modified differential display of tumour antigens, in the method of the invention. The specification describes a method for identifying a target epitope. The method comprises screening the products of an expression library from a cell expressing the target epitope with cytotoxic T cells generated
Sequence 10 BP; 3 A; 3 C; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel method of identifying target epitopes or antigens specific for human tumors, cancers and infected cells involving screening express: library products of a cell expressing the target epitope.
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G; 3 T; 0 U; 0 Other;
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RESULT 58
AAF35009/c
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                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence is a primer used to detect polymorphisms in the human osteoclastogenesis inhibitory factor (TNFRSF11B). Polymucleotides comprising one or more of twenty four novel single nucleotide polymorphisms in the TNFRSF11B gene have been identified. TNFRSF11B regulate osteoclast recruitment and function. An understanding of variations in the gene should thus be useful in developing new therapies for metabolic disorders caused by abnormal osteoclast recruitment and function such as osteoporosis, metastatic bone disease, Paget's disease, rheumatoid arthritis and periodontal bone disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; TNFRSF11B; osteoclastogenesis inhibitory factor; single nucleotide polymorphism; SNP; osteoclast recruitment; osteoclast function; osteoporosis; metastatic bone disease; Paget's disease; rheumatoid arthritis; periodontal bone disease;
                                                                                        nor previously assigned open reading serial analysis of gene expression; a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human Osteoclastogenesis Inhibitory Factor nucleotides, comprising single nucleotide polymorphisms, useful for studying e.g. osteoporosis, Paget's disease and rheumatoid arthritis.
                                                  Saccharomyces cerevisiae
                                                                                                                                                 Yeast NORF
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        primer extension; primer; ss.
21-DEC-2000
                                                                                                                   Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;
                                                                                                                                                                          23-MAR-2001
                                                                                                                                                                                                                                AAF35009
                                                                                                                                                                                                                                                                                                                          1496 ACTTCAGCA 1504
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                                                                                                                                                                                                                                                                                                                                                    Similarity 9; Conserv
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                                                                                                                                                                                                                                standard; DNA; 10
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                                                                                                                                                gene
                                                                                                                                                                                                                                                                                                                                                                                                         BP; 2 A;
                                                                             primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page
                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                          (first entry)
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                                                                                                                                                SAGE tag oligonucleotide SEQ ID NO:1748.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24;
                                                                                                                                                                                                                                                                                                                                                                                                         1 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      114pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Duda A, Nandabalan K, Stephens JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer-extension oligo,
                                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                                                                                                                               30.0%;
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                                                                                                                                                                                                                                                                                                                                                                               Score 9;
                                                                                                                                                                                                                                                                                                                                                                   Pred.
                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                   No.
                                                                                          g frame; nonannotated
antifungal; tag; ider
                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ
                                                                                                                                                                                                                                                                                                                                                                             Length 10;
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RESULT 59
AAF39472/c
ID AAF394
XX AAF394
XX AAF394
XX Yeast
XX Yeast
XX Yeast
XX Yeast
XX Yeast
XX Yeast
XX Inner px
XW Serial
XW Serial
XX Saccha
XX Saccha

Yeast NORF

gene

SAGE tag oligonucleotide

SEQ ID NO:6211

23-MAR-2001

(first entry)

AAF39472 standard; DNA; 10

ВÞ

Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;

nor previously assigned open reading serial analysis of gene expression; linker; PCR primer; ds.

밁 S

10

TTCAGCAGC

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Matches Query Match

Similarity 9; Conserv

Conservative

30.0%;

Score 9; Pred. No

<u>ب</u>

Length 10

Mismatches No. DB .

0

0

Gaps

0

1498 TTCAGCAGC 1506

Sequence 10 BP; 2 A; 2 C; 3 G; 3 T; 0 U; 0 Other;

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cc antifugal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human pares which are involved in cell cycle progression of comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; candidate of the control of the present invention.

Cc pares of the control of the present invention of method, in the exemplification of the present invention of method, in the exemplification of the present invention of the pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonamnotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a year
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-JUN-2000; 2000WO-US016223
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             62; 419pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9908-00335032
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RESULT 60
AAF38820
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     Query Match
Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell of the cell cycle, the differentially expression which will be a control of the cell cycle, the differentially expression where the control of the cell cycle, the differentially expression where the control of the cell cycle, the differential
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF33268 to AAF4064 represent SAGE tags used in the exemplification of the present invention AAF33362 to AAF33267 represent linkers and PCR primers used in the SAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes an isolated DNA molecule comprising coding sequence of a yeast gene selected from a group of 745 NORF (repreviously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                                                                                                                                                                                                                         AAF38820 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 10 BP; 1 A; 2 C; 2 G; 5 T; 0 U; 0 Other;
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                                                                                                                                         Yeast
                                                                                                                                                                                                23-MAR-2001
                                                                                                                                                                                                                                                     AAF38820
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1483 ACCAAGAAG 1491
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                                                                                                                                      gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                             (first entry)
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                                                                                                                                         SAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                   tag oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           score 9; DB 1; Pred. No. 43; 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
o. 43;
                                                                                                                                         SEQ
                                                                                                                                         IJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 10;
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                                                                                                                                         NO:5559
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AAF41980
ID AAF4
XX AC AAF4
XX AC AAF4
XX Yeas
XX Yeas
XX Yeas

23-MAR-2001

(first entry)

AAF41980;

AAF41980 standard;

DNA;

10

Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;

Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:8719

밁 5

Matches

9;

Conservative

0

Mismatches

Indels

٥,

0

1486 AAGAAGCCA 1494

AAGAAGCCA 9

Query Match Best Local Similarity

30.0%;

Score 9; ; Pred. No.

DB 1; o. 43;

Length 10; ٥,

RESULT 61

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Cycle comprising administering a NORF gene whose expression varies by at Ciphase, S phase and G2/M; (2) a method (M2) for screening candidate CC antifungal drugs comprising: (a) contacting a test substance with a yeast Cell; and (b) monitoring expression of a NORF gene whose expression of CC varies as in M1, where a test substance which modifies the expression of CC the yeast gene is a candidate antifungal drug; (3) a method (M3) for CC identifying human genes which are involved in cell cycle progression CC comprising contacting human DNA with a probe which comprises at least 10 CC contiguous mucleotides of a NORF gene whose expression varies as in M1; CC and (4) a method (M4) for identifying a candidate drug as a member of a CC class of drugs having a characteristic effect on gene expression in a CC class of drugs having contacting a yeast cell with a candidate drug and CC expression is affected by the class of drugs. The NORF gene whose expression is affected by the class of the cell cycle, the differentially CC expressed genes may be used as markers of phases of the cell cycle. The CC cycle and for identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF33268 to AAF44064 CC cycle and for identification of the present invention. CC AAF33267 to AAF3267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saccharomyces cerevisiae.
Sequence 10 BP; 6 A; 2 C; 2 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          affecting phases of the cell cycle.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example; Page 198; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-061874/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Velculescu V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9908-00335032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kinzler
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                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene sexpression) tag. Also comprising a SAGE (sexial analysis of gene expression) tag. Also code comprising a sade (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at cell east 10% between any two phases of the cell cycle selected from log code antifungal drugs comprising; (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of codentifying human genes which are involved in cell cycle progression code the yeast gene is a candidate antifungal drug; (3) a method (M3) for contiguous nucleotides of a NORF gene whose expression of codentifying a method (M4) for identifying a candidate drug as a member of a constraining contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell with a candidate drug and code cexpression is affected by the class of drugs. The NORF gene whose expression in a control of expression in the yeast cell of at least 1 NORF gene whose cexpression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The cycle and for identification of antifungal drugs which affect the cell cycle. The method in the same time of the present invention.

Comprise and same of the creamplification of the present invention. The method in the SAGE translation of the present invention.
XXXXXX
                                                                                                            AAF38821
                                                                                                                                RESULT 62
                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                23-MAR-2001
                                                      AAF38821;
                                                                                                                                                                                                                                                                                                                                            Sequence 10
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                                                                                        AAF38821 standard; DNA; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-JUN-2000; 2000WO-US016223
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              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                 exemplification of the present invention
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                                                                                                                                                                                                                                                                                  30.0%;
                                                                                                                                                                                                                                                                                                                                            2 C; 4
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                                                                                                                                                                                                                                                                                    Score 9;
Pred. No.
                                                                                                                                                                                                                                                                                                                                            G; 0 T; 0
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                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                            U; 0 Other;
                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                      43;
                                                                                                                                                                                                                                                                                                      Length 10;
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Yeast NORF gene SAGE tag
                                                 oligonucleotide
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                                                 NO:5560
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Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.

Saccharomyces cerevisiae

14-JUN-2000; 2000WO-US016223

16-JUN-1999; 9975-00335032

(UYJO) UNIV JOHNS HOPKINS

Velculescu V, Vogelstein B, Kinzler

affecting phases of the cell cycle. Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and of.

Example; Page 198; 419pp; English.

The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not comprising assigned open reading frame; or nonannotated ORF) genes comprising assigned open reading frame; or nonannotated ORF) genes comprising alministering a NORF gene expression tag. Also cycle comprising administering a NORF gene whose expression varies by at certain 10% between any two phases of the cell cycle selected from log chase, S phase and G2/M; (2) a method (M2) for screening candidate cell; and (b) monitoring expression of a NORF gene whose expression of cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for contiguous nucleotides of a NORF gene which modifies the expression of comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The Corp. and for identification of antifungal drugs which affect the cell cycle and for identification of antifungal drugs which affect the cell cycle. The present shows the present invention.

CC method in the seemalification of antifungal drugs and CR primers used in the SAGE method in the SAGE in in the exemplification of the present invention

Sequence 10 B₽; 5 A; 2 C; 2 G; 1 T; 0 U; 0 Other;

밁 S Query Match Best Local S Matches 9 1486 _ Similarity 9; Conserv AAGAAGCCA 1494 Conservative 100.0%; 30.0%; 0 Score 9; 1 Pred. No. Mismatches DB 1 ۳. Length 10; Indels 0

Gaps

0

RESULT 63
AAF39528
ID AAF39
XX
AC AAF39 AAF39528 AAF39528 standard; DNA; 10

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S
                                                                                                                                                                cc antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for contacting human genes which modifies the expression of cc dentifying human genes which are involved in cell cycle progression cc comprising contacting human DNA with a probe which comprises at least 10 cc contiquous nucleotides of a NORF gene whose expression varies as in M1; cand (4) a method (M4) for identifying a candidate drug as a member of a cc class of drugs having a characteristic effect on gene expression in a cyeast cell comprising contacting a yeast cell with a candidate drug and cc waste cell comprising contacting a yeast cell with a candidate drug and cc expression is affected by the class of drugs. The NORF gene whose cexpression is affected by the class of the cell cycle, the differentially contacted genes may be used as markers of phases of the cell cycle. The cethods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungs. AAF33268 to AAF44064 cc represent SACE tags used in the exemplification of the present invention. Cc AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE contaction of the present invention.
                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF gene affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate
                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes an isolated DNA molecule comprising coding sequence of a yeast gene selected from a group of 745 NORF ()
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
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                                                                                                                                                              BP; 4 A;
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                                                                                                                                                            2 C; 3 G; 1 T; 0 U; 0 Other;
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Pred. No.
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RESULT 64 AAF38557

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1489

AAGCCAGAC 1497 AAGCCAGAC

10

Query Match Best Local S Matches 9

Similarity 9; Conserv

Conservative

0

30.0%; Score 9; 1 100.0%; Pred. No.

DB 1;

Length 10

0

Gaps

0;

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cc antifungal drugs comprising: (a) contacting a test substance with a yeast cell, and (b) monitoring expression of a NORF gene whose expression of varies as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for cidentifying human genes which are involved in cell cycle progression of comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a cyeast cell comprises in contacting a yeast cell with a candidate drug and contacting expression in the yeast cell with a candidate drug and conticring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of the cell cycle, the differentially contacting and feet phases of the cell cycle, the differentially contact may be used as markers of phases of the cell cycle. The contact may be used as markers of phases of the cell cycle. The contact may be used as markers of phases of the cell cycle. The cycle and for identification of antifungal drugs. AAF33268 to AAF34064 crepresent SAGE tags used in the exemplification of the present invention.

CC method, in the exemplification of the present invention
Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       previously assigned open reading frame; or nonannotated ORF) genes comprising a SASE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonamotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example; Page 189; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:5296
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BP; 5 A; .3 C; 2
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G; 0 T; 0 U; 0 Other;
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RESULT 65
AAD31785
                                                     Query Match
Best Local Similarity
'---hes 9; Conserve
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                                                                                                                                                                                                                                                                                            antigens recognised by cytotoxic T cells (CTLs) and specific for human tumours, cancers and infected cells. The method involves screening the products of an expression library generated from DNA/RNA of a cell products of an expression library generated from DNA/RNA of a cell capressing a target epitope with cytotoxic T cells generated against the cytotoxic T cells specific for a gene product differentially expressed by capressing the cross-reactivity of the cytotoxic T cells for cells for cells expressing a target epitope in which the target epitope is dentified as a gene product inducing cytotoxic T cells. The method is useful for identifying a target epitope or antigen specific for a tumour cell. The target epitope is also useful for identifying target epitope or antigen specific for a fumour cell. The target cells against which it is desirable to induce cell mediated immunity. The antigen identified by the method is useful in capression of tumours, cancers and infections in mammals. The invention capression for treating to the relates to vaccinia viral vectors which are useful for treating the resonance.
                                                                                                                                                                                 tumour-bearing mammals, including humans to generate immune response against the tumour cells. They are also useful for immunising or vaccinating tumour-free such tensor tumour formation. The present DNA sequence is an arbitrary primer which is used for modified differential display of genes encoding potential tumour immunogens. This primer is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Representational Difference Analysis method for identification of antigens recognized by cytotoxic T cells and specific for human tumors,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD31785;
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                                                                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 4; Page 19; 54pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   comprises improved selection of genes encoding target antigens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-APR-2001; 2001US-00822250
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                                 1493 CAGACTTCA 1501
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CAGACTTCA 10
                                                                                                                                                BP; 3 A; 3 C; 1 G; 3 T; 0 U; 0 Other;
                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l; CTL; tumour; cancer; infection; cell-mediated immunity;
response; cytostatic; primer; ss.
                                                                                        30.0%;
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                                                                                      Score 9; I
                                                                       Mismatches
                                                                                      DB 1;
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                                                                       Indels
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RESULT 66 ABQ87208/c

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RESULT 67
ABV69313/c
ID ABV693
XX
AC ABV693
AC ABV693
XY
DT 21-OCT
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DE Human
XX
DT Human;
KW immuno
KW jeoria
XX
DS Homo s
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PN WO2002
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        expression. (A) comprises protein or mRNAs or their fragments. (M1) is useful for: identifying markers of skin ageing and/or stress; determining skin ageing and/or stress; and identifying or determining the effects of pharmaceutical or cosmetic agents for control of skin ageing. The present sequence is one of a group of human skin ageing/stress related expressed sequence tags (ABQ86246-ABQ87680) of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to identifying (M1) genes in vitro that, in humans or animals, are important for skin ageing and/or skin stress by serial analysis of gene expression between mixtures of transcribed and optionally translated, genetically encoded factors (A) obtained from young and aged skin, to identify that genes that show strong differential expression. (A) comprises protein or mRNAs or their fragments. (M1) is
                                                                        Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                       Human skin
                                                                                                                                                                     21-OCT-2002
                                                                                                                                                                                                  ABV69313;
                                                                                                                                                                                                                               ABV69313 standard; cDNA; 11
                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 11 BP; 1 A; 2 C; 4 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Identifying genes involved in skin stress and aging, useful e.g. in screening for cosmetic or therapeutic agents, based on differential gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-528865/56.
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                                             Homo sapiens.
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                                                                                                                                       EST
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                 BG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                              WPI; 2002-590638/63
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In vitro identification of skin-expressed genes, useful for determining
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                     The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin
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   disorders, specifically neurodermatitis; sunburn;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human skin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABV66240;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HENK ) HENKEL KGAA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABV66240 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1490
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9; Conserv
                                                                   9
TTCAGCAGC 1506
                                                                                                                                                                                                    the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGCCAGACT 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGCCAGACT 1498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EST
                                                                                                                                                       BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 136; 1345pp; German.
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                       3 A;
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100.0%; Pr
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                                                                                                                                                       3 C; 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hofmann
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Pred. No.
                                                                                     Score 9; I
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                                                                     Mismatches
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                                                                                                                                                       T; 0 U; 0 Other;
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                                                                                     DB 1;
5. 47;
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). 47;
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                                                                                                            Length 11;
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RESULT 71
ABV67707/c
ID ABV67707/s
XX ABV67707;
AC ABV677
RESULT 72
AAQ57385/c
ID AAQ573
XX
AC AAQ573
XX
AC AAQ573
C
DT 25-MAR
DT 26-JUL
XX
DE Enzyma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     in the skin of humans or animals by subjecting a mixture of genes expressed encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes and quantify their expression determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitie.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
Enzymatic RNA
                                                         25-MAR-2003
26-JUL-1994
                                                                                                                                               AAQ57385;
                                                                                                                                                                                                         AAQ57385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 11 BP; 1 A; 3 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disorders, specifically neurodermatitis; sunburn; psoriasis; sclerodermatichthyosis; atopic cermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the second of the second part of the present sequence is that of a human expressed sequence tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 176; 1345pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            e.g. skin cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-590638/63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-JAN-2001; 2001DE-01000127
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                                                                                                                                                                                                                                                                                                                                                                                                                  1500 CAGCAGCCA 1508
                                                                                                                                                                                                                                                                                                                                                           11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      l Similarity
9; Conserva
                                                                                                                                                                                                         standard; mRNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention
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                                                         (revised)
(first entry)
molecule ACE mRNA target sequence.
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o. 47;
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Best Local S
Matches 9
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07-DEC-1992;
07-DEC-1992;
07-DEC-1992;
19-JAN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This is a ACE mRNA target sequence (nucleotide no. 2076) of an enzymation and coule (ribozyme) which cleaves mRNA associated with the development or maintenance of a cardiovascular condition. The concn. of the ribozyme necessary to effect a therapeutic treatment is lower than that of an antisense oligonucleotide and the specificity of action is higher. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide primer SEQ ID NO 355765 for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Specific; cleavage; target RNA; protein; prophylaxis; expression; inhibitor; inhibition; ribozyme; treatment; prevention; psoriasis; asthma; inflammatory diseases; cardiovascular condition, hypertension; arthritis; restenosis; angiotensin converting enzyme; ss.
07-APR-2000; 2000DE-01019173
                                                     06-APR-2001; 2001WO-IB000713
                                                                                                                                                                                                                                                                                                                                                                                                                                       22-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABI55792 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 3; Page 23; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Enzymatic RNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inflammatory,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1500 CAGCAGCCA 1508
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CAGCAGCCA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12 BP; 1 A; 3 C; 5 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      molecules which cleave \mathfrak{m}RNA - used to treat or prevent arthritic, stenotic or cardiovascular diseases or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Draper KG
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92US-00987132.
92US-00989848.
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93US-00008895
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 T; 0 U; 0 Other;
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5. 51;
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                                                                                                                                                                                                                                                                                                                                                                                    SNP TSC0049803.
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                                                                                                                                                                                                                                                                           metabolic.
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RESULT 74
ABH9393
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Best Local S
Matches 9
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                                                             Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
                                                                                                                                                                                                                                                                                   07-APR-2000; 2000DE-01019173
                                                                                                                                                                                                                                                                                                                           06-APR-2001; 2001WO-IB000713.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        designed to detect methylation status.
                                                                                                                                                 WPI; 2001-657177/75
                                                                                                                                                                                              Olek
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                                                                                                                                                                                            Piepenbrock C,
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                            range of diseases including immune system, gastrointestinal, respiratory central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation, ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>,,</u>
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                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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RESULT 78
AAQ99836
ID AAQ99
XX
AC AAQ99
XY
COMMAND
COM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010—ABC9989, ABF00010—ABC9989, ABF00010—ABC998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            set or oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
       Synthetic
                                                               Eucalyptus; urophylla; grandis; wood volume marker; RAPD genetic marker; random amplified polymorphic DNA analysis; woody perennial plant; family selection; pedigree; mapping; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 12 BP; 5 A; 4 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                  Eucalyptus
                                                                                                                                                                                                                                                                                                                 06-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                         AAQ99836;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ99836 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 377549; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-657177/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01ek
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-APR-2001; 2001WO-IB000713
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200177384-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide primer SEQ ID NO 377549 for detecting SNP TSC0062385.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (EPIG-) EPIGENOMICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABI77576 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1479 CACGACCAA 1487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ۶
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CACGACCAA 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Piepenbrock C,
                                                                                                                                                                                                                                      wood volume
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                cDNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30.0%;
                                                                                                                                                                                                                                  marker half-sib primer V7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Berlin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 9; DB 1; Pred. No. 51; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>~</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
o. 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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RESULT 79
AAV50097
ID AAV50
XX AAV50
XX Yeast
XX Yeast
XX Yeast
XX Yeast
XX Sacch
OS Synth
XX WO983
XX Sacch
PD 30-JI
XX Z-JJ
XX Z-JJ
XX WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Watches 9; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yeast; Saccharomyces cerevisiae; transcriptome; cell cycle; eukaryotic cell; antifungal; SAGE tag; gene expression; serial analysis of gene expression; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Determn. of heritable oligogenic traits in woody plants by genomic mapping of multiple markers in a two generation plant family - uses select plants with desired characteristics for breeding.
Yeast transcriptome - useful for screening antifungal agents, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 10 BP; 3 A; 4 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Omalley
                                           WPI; 1998-427943/36.
                                                                                                                                                                               30-JUL-1998
                                                                                                                                                                                                          WO9832847-A2
                                                                                                                                                                                                                                                 Saccharomyces
                                                                                                                                                                                                                                                                                                                                  Yeast tag
                                                                                                                                                                                                                                                                                                                                                              21-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                   AAV50097 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          fragments from the genomic DNA of E.urophylla, E.grandis and F1 probtained by crossing the two species. Subsequent mapping analysis that the half-sib primers in AAQ99834-Q99840 are all useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RAPD analysis was used to determine whether certain quantitative traits were heritable oligogenic traits in Eucalyptus trees. Sets of commercially available random 10-mer primers were used to amplify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 6; Page 58; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1995-269212/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-JAN-1995;
                                                                      Velculescu
                                                                                                (UYJO ) UNIV JOHNS HOPKINS SCHOOL MEDICINE.
                                                                                                                           23-JAN-1997;
                                                                                                                                                    22-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amplifying wood volume markers from Eucalyptus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYNC-) UNIV NORTH CAROLINA STATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-JAN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9519697-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1488 GAAGCCAGAC 1497
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GAAGCCAGCC 10
                                                                                                                                                                                                                                                                                                                                for putative coding sequence locus YBR162C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sederoff
                                                                                                                                                                                                                                                  cerevisiae
                                                                                                                           97US-0035917P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95WO-US000677.
                                                                                                                                                    98WO-US001216
                                                                   Vogelstein
                                                                                                                                                                                                                                                                                                                                                                                                                     DNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RR,
                                                                     B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Grattapaglia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 8.4; I
Pred. No. 56;
                                                                      Kinzler KW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
modulating eukaryotic cell, for for identifying genes in cell cycle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ā
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DΒ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                        regulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ٥,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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RESULT 80
AAX59809
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       an isolated DNA molecule comprising a yeast gene which is involved in cell cycle progression selected from differentially expressed genes (SAGE tags given in AAVS0051 to AAVS0345); (2) a method for screening candidate antifungal drugs which comprises contacting a test substance with a yeast cell and monitoring expression of a yeast gene which is involved in cell cycle progression; (3) a method of identifying human genes which are involved in cell cycle progression which comprises hybridizing a probe comprising at least 10 contiguous nucleotides of a yeast gene which is differentially expressed between at least 2 phases selected from the log phase, the S phase and the GCZ/M phase; and (4) a probe for ascertaining the phase in the cell cycle, where the probe comprises at least 14 contiguous nucleotides of a NORF gene (SAGE tags given in AAVS0051 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yeast transcriptome is encoded by a DNA molecule comprising a yeast gene involved in cell cycle progression selected from the group of nonannotated ORF (NORF) genes. SAGE (serial analysis gene expression) tags for highly expressed genes and NORF genes are given in AAV50051 to AAV50345. The present invention describes: (1) a method of using yeast genes to modulate the cell cycle which comprises administering to a cell genes to modulate the cell cycle which comprises the comprise of the cell cycle which comprises administering to a cell
                                                                                                                                                               Grattapaglia
Kuhlman EG;
                                                                                                                                                                                                                                                                                                                                                                                            Genetic marker; genetic locus; resistance; fusiform rust disease; tree family; Pinus; volume marker; PCR primer; ss.
                                                                                                                                                                                                                                                                                                       01-JUN-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Primer V7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX59809
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX59809 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 10 BP; 4 A;
                                                        Example
                                                                                                                                    WPI; 1999-347038/29
                                                                                                                                                                                                                                           21-JAN-1994;
                                                                                                                                                                                                                                                                       18-OCT-1995;
                                                                                                                                                                                                              (UYNC-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1480 ACGACCAAGA 1489
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 9; Conserv
                                                        6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACGGCCAAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 22; 44pp; English
                                                          6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or as an array of probes on a solid support
                                                                                                      resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                               'n
                                                                                                                                                                                                             NORTH CAROLINA STATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                        amplifying volume markers in plant half-siblings.
                                                       37; 69pp;
                                                                                                                                                                                                                                           94US-00184567.
                                                                                                                                                                                                                                                                       95US-00545253
                                                                                                                                                                             O'malley DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 C; 3
                                                                                                      to fusiform
                                                        English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 8.4; D
Pred. No. 56;
                                                                                                                                                                                Amerson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                      rust disease
                                                                                                                                                                               ,VH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DΒ
                                                                                                                                                                                Sederoff
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Length 10;
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                                                                                                      'n
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                                                                                                      trees
                                                                                                                                                                                RR,
                                                                                                      of the
                                                                                                                                                                                Wilcox
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                                                                                                      genus
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The specification describes a method of identifying a genetic marker associated with a genetic locus conferring resistance to fusiform rust disease in a family of trees of the genus Pinus. The method comprises

Sequence

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0 ς,

0 Other;

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RESULT 81
AAZ22666
ID AAZ22666
ID AAZ22666
ID AAZ22
XX AC AAZ22
XX DT 04-JJ
XX COMMME
KW Prime
XX Synth
XX EP950
XX EP950
XX 16-MJ
XX 16-M
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        obtaining a sexually mature Pinus parent tree exhibiting resistance to fusiform rust disease, obtaining a plurality of progeny trees of the parent by self or cross-pollinations, assessing multiple progeny trees for a number of genetic markers, identifying genetic markers segregating in a Mendelian ratio and showing linkage with other genetic markers measuring resistance to fusiform rust disease in multiple progeny trees and correlating the presence of resistance to fusiform rust disease with at least one marker identified in the previous step. The method is useful for determining the genetic basis of resistance to fusiform rust disease and for producing trees of the Pinus genus that are resistant to the disease. The present primer was to amplify volume markers in plant half-siblings, in the course of invention
                                                                                  This is a commercial oligonucleotide primer for the PCR-based RAPD analysis of fungal isolates of Dydimella bryiniae and Phoma species. The new method may be used to distinguish D. bryoniae from non-pathogenic Phoma species fungal infections in Cucurbits (i.e.cucumbers, pumpkins, watermelons, gourds, cantaloupes, squashes and related plants). The new method of detection of D. bryoniae and Phoma species infections allows rapid diagnosis even before symptoms are visible as compared to prior art methods which involved growing pure cultures of the pathogens from the infected plants and identifying them under the light microscope. The method leads to the early treatment of the infected plants with fungicides resulting in an increased chance of saving the infected food
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-MAR-1998;
22-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EP950719-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           commercial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    T18 primer for amplification of fungal genomic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-JAN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 10 BP; 3 A; 4 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           recombinant techniques.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detecting a pathogenic fungus in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-582557/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-MAR-1999;
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nes 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GAAGCCAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        assay; test; fungal pathogen;
; RAPD analysis; isolate; food
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Somai BM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLEMSON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0078103P
99US-00255432
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              8; 1.9pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              English.
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Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cucumbers,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            crop protection; cucurbit;
crop; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pumpkins and gourds using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 10;
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Best Local Similarity
Matches 9; Conserv
                                                                                                                              19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   SAGE
APC;
Isolated polynucleotides differentially expressed cells, useful in gene vaccines against cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human dendritic cell SAGE tag,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ79223 standard; DNA; 10 BP.
                          WPI; 2000-106077/09.
                                                           (GENZ )
(ROBE/)
(SHAN/)
                                                                                                                                                                                                                                                                        19-JUN-1998
                                                                                                                                                                                                                                                                                                                                              19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                 cytotoxic T-lymphocyte; tumour antigen; immunotherapy; anticancer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-APR-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ79223;
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                                                                                                                                                                                                                                                                                          19-JUN-1998
                                                                                                                                                                                                                                                                                                   19-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                       AGE tag; serial analysis of gene expression; antigen-presenting cell; APC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1488 GAAGCCAGAC 1497
                                                                                                                                                                                    1-1998
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                                                           GENZYME CORI
ROBERTS B L.
SHANKARA S.
                                          BL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GATGCCAGAC 10
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                                          Shankara
                                                                                                                              98US-0090036P

98US-0090040P

98US-0090041P

98US-0090042P

98US-0090044P

98US-0090044P

98US-0090044P

98US-0090047P

98US-0090047P

98US-0090076P

98US-0090076P
                                                                                             98US-0090078P.
98US-0090079P.
98US-0090080P.
98US-0111715P.
                                                                                                                                                                                                                                               98US-0090000P.
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                                                                   CORP.
                                                                                                                                                                                                                                                                        98US-0089994P.
98US-0089997P.
                                                                                                                                                                                                                                                                                         98US-0089993P
                                                                                                                                                                                                                                                                                                          98US-0089878P
98US-0089991P
                                                                                                                                                                                                                                                                                                   98US-0089992P
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Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO:1651
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      56;
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        antigen-presenting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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SX Claim 1; Page 112; 130pp; English.

XX Sequences AAZ77573-Z79709 represent SAGE (serial analysis of gene contents of expression) tags used to identify mRNA transcripts encoding climmunostimulatory cofactor proteins which are preferentially or differentially expressed in monocyte-derived dendritic cells compared contents of the transcripts correspond to known genes or ESTs (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while cother transcripts correspond to novel genes. Antigen-presenting cell (APC)-associated costinulatory factors play an important role in the cytotoxic immune response, particularly against tumour cells. Tumour antigen presentation via the MHC (major histocompatibility complex) and subsequent recepnition by T-cell receptors is alone cefficient activation of tyetoxic T-lymphocytes (CTas). Nucleic acid cefficient activation of cytotoxic T-lymphocytes (CTas) nucleic acid cefficient activation of cytotoxic T-lymphocytes (CTas). Nucleic acid can APC; and as hybridisation probes/amplification primers for the cappression of these genes to induce an immune response, particularly capainst a tumour antigen; to modulate the genotype of an APC; to screen cefficient activation of these genes of their encoded proceins, can be used to identify ceptures of the monocyte illneage. Cells containing these genes in the cappression of these genes. Detection of the dendritic cell differentially capressed genes, or of their encoded proceins, can be used to identify immunotherapy (or costimulate production of a complex costimulatory factors ensures adequate antigen and cetors and secontaining these genes cappulation of and vectors containing these genes cappulation of cell-growth factors and secretion of cell-rich sites, servention of T cell growth factors and secretion of chemokines for the presentation of cell-rich sites, sequence 10 BP; 3 A; 4 C; 3 G; 0 T; 0 U; 0 Other;
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RESULT 83
AAZ82715
ID AAZ82
XX
AC AAZ8.
DT 07-A
XX
DE Meta
XX
CHARACTER MATE
XX
DE MOTO
XX
HUMA
KW HUMA
KW AND
XX
PN WOS
XX
PN WOS
XX
PP 18
XX
PR 15
PR 15
PR 115
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Best Local Similarity
Matches 9; Conserv
 19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss.
                                                                                                                                                                                                                    Metastatic breast tumour cell upregulated transcript tag #1949.
                                                                                                                                                                                                                                               07-APR-2000
                                                                                                                                                                                                                                                                       AAZ82715;
                                                                                                                                                                                                                                                                                               AAZ82715 standard; DNA; 10
                                                                                         23-DEC-1999
                                                                                                                WO9965928-A2
                                                                18-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                        1488 GAAGCCAGAC 1497
                                                                                                                                         sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                             (first entry)
             98US-0089853P.
98US-0089997P.
98US-0090039P.
                                                                99WO-US013647.
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90.0%;
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Pred. No. 56;
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RESULT 84
AAZ81205/c
ID AAZ812
XX AAZ812
XX AAZ812
XX O7-APR
XX Metast
XX Human;
KW non-me
KW antime
KW antime
KW antime
XX Homo s
XX Homo s
XX Homo s
XX Homo s
XX 19-JUN
PR 19-JUN
PR 19-JUN
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Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for adoptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942
                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                      Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer;
                                                                                                                                                                                                                                                                                                               07-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 10 BP; 4 A; 4 C; 1 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 111; 219pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Roberts BL,
   19-JUN-1998;
19-JUN-1998;
                                                        18-JUN-1999;
                                                                                                                                WO9965928-A2
                                                                                                                                                                                                                                                                                                                                                   AAZ81205
                                                                                                                                                                                                                                                                                                                                                                                    AAZ81205 standard; DNA; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENZYME CORP. ROBERTS B L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SHANKARA S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TCAACAGCCA 10
                                                                                                                                                                                                                                                                             breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                               (first entry)
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   98US-0089853P.
98US-0089997P.
                                                        99WO-US013647.
                                                                                                                                                                                                                                                                       tumour cell upregulated transcript tag #439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28.0%;
90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ß
                                                                                                                                                                                                                  tumour tissue; gene therapy;
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Pred. No. 56;
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RESULT 85
AAZ82619
ID AAZ82
XX AAZ82
XX AAZ82
XX O7-AF
DT O7-AF
XX Human
KW Human
KW INON-H
KW ANTIH
OS HOMO
XX WO996
XX WO996
XX WO996
XX WO996
XX IB-JI

antimetastatic;

non-metastatic breast tumour tissue; gene

vaccine; diagnosis;

Human; metastatic breast tumour tissue; breast cancer; tag;

therapy; anticancer;

18-JUN-1999; 23-DEC-1999

99WO-US013647.

WO9965928-A2

Metastatic breast tumour cell upregulated transcript tag #1853

07-APR-2000

(first entry)

AAZ82619 standard; DNA; 10

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1486 AAGAAGCCAG 1495

AAGAAGGCAG

Query Match Best Local (Matches

Similarity 9; Conserv

Conservative

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Mismatches

1;

0,

Gaps

0

28.0%;

Score 8.4; I Pred. No. 56;

DB 1;

Length 10;

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CC that are preferentially transcribed in the metastatic breast tumour colls, are preferentially transcribed in the metastatic breast tumour cells). AAZ83942 CC tissue (i.e. are upregulated in metastatic breast tumour cells) to AAZ866677 represent tags corresponding to distinct transcripts that are cc preferentially transcribed in the primary or non-metastatic breast tumour cells). These crises (i.e. are downregulated in metastatic breast tumour cells). These crises (i.e. are downregulated in metastatic breast tumour cells). These crises (i.e. are downregulated in metastatic breast tumour cells). These crises (i.e. are downregulated in metastatic breast tumour cells). These crises (i.e. are downregulated in metastatic) breast tumour cells). These crises (i.e. are downregulated in metastatic) breast tumour cells. Compounds that modulate expression of the transcripts are potentially cuseful for treatment of (metastatic) breast cancer, while promoters from crises for treatment of (metastatic) breast cancer, while promoters from crises promoters from crises promoters are used to direct expression, in selected cell types, of ce.g. therapeutic genes (also ribozymes or antisense sequences), of crises, for diagnosing breast cancer and for raising specific cagnits. Host cells that produce the polypeptides can be useful in cells, especific and isolate populations of educated, antigen-specific immune effecter community effecter.
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19-JUN-1998;
19-JUN-1998;
Sequence 10 BP; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-106079/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            treatment of cancer.
                                         immunotherapy
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ROBERTS B L.
SHANKARA S.
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98US-0090040P.
98US-0090041P.
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A; 4 C; 1
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<u>Ω</u>
5 T; 0 U; 0 Other;
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AAZ84503
ID AAZE
XX
AC AAZE
XX
AC AAZE
DT 07-J
XX
Meta
XX
Meta
XX
Humm
KW non.
KW anct
XX
Hom
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts CC that are preferentially transcribed in the metastatic breast tumour CC tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 CC to AAZ86677 represent tags corresponding to distinct transcripts that are CC preferentially transcribed in the primary or non-metastatic breast tumour ctissue (i.e. are downregulated in metastatic breast tumour cells). These CC transcripts can be used for diagnosis, prognosis, monitoring and CC treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. CC Compounds that modulate expression of the transcripts are potentially CC useful for treatment of (metastatic) breast cancer, while promoters from CC the transcripts are used to direct expression, in selected cell types, of cell. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based CC vaccines. Polypeptides encoded by the transcripts are also useful in CC vaccines, for diagnosing breast cancer and for ratsing specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic capits. Host cells that produce the polypeptides or as therapeutic colls, e.g. cytotoxic T lymphocytes, and these used for adoptive
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Matches
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19-JUN-1998;
19-JUN-1998;
                               23-DEC-1999
                                                                                                                                                                                                                                          Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer;
                                                                                                                                                                                                                                                                                                                                  Metastatic breast tumour cell downregulated transcript tag
                                                                                                                                                                                                                                                                                                                                                                                                 07-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ84503 standard;
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                                                                                              WO9965928-A2
                                                                                                                                                                                                                   antimetastatic;
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(ROBE/)
(SHAN/)
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) ROBERTS B L.
) SHANKARA S.
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9; Conserv
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98US-0089997P.
98US-0090039P.
98US-0090040P.
98US-0090041P.
                                                                                                                                                                                                                   vaccine; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA; 10
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RESULT 87
AAZ84255
ID AAZ84
XX
AC AAZ84
XX
DT 07-Ai
DT 07-Ai
XX
Humar
KW Humar
KW mon-r
KW antir
XX
XX
Homo
OS Homo
XX
XX
WO99
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Best Local S
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These
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19-JUN-1998;
19-JUN-1998;
                                                                                                                 Human; metastatic breast tumour tissue; breast cancer; tag; primer;
                                                                                                                                              Metastatic breast tumour cell downregulated transcript tag #3489.
                                                                                                                                                                                07-APR-2000
                                                                                                                                                                                                               AAZ84255;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 10 BP; 6 A; 1 C; 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GENZ )
(ROBE/)
(SHAN/)
                                                                                                non-metastatic
                                                                                                                                                                                                                                                AAZ84255 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              non-metastatic breast cancer cells, useful for diagnosis, treatment of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-JUN-1999;
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) ROBERTS B L.
) SHANKARA S.
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9; Conserv
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98US-0089997P.
98US-0090039P.
                                                                                              breast tumour tissue; gene therapy; anticancer;
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                                                                                                                                                                                                                                              DNA;
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Pred. No.
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Homo sapiens

W09965928-A2

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RESULT 88
AAZ81942/c
ID AAZ819
XX AAZ819
AC AAZ819
XX O7-APR
XX DE Metast.
XX Human;
XW non-me
XW antime
XX AX8
OS Homo s
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                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 9; Conserv
                                                  Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss.
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(ROBE/)
                                                                                                                          Metastatic
                                                                                                                                                           07-APR-2000
                                                                                                                                                                                                                             AAZ81942 standard; DNA; 10 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Isolated polynucleotides differentially expressed between metastatic non-metastatic breast cancer cells, useful for diagnosis, prevention
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ROBERTS B L.
SHANKARA S.
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98US-0089997P.
98US-0090039P.
98US-0090040P.
98US-0090041P.
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                                                                                                                       tumour cell upregulated transcript
                                                                                                                                                                                                                                                                                                                                                                                                   28.0%;
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AAA11232
ID AAA11
XX AAA11
XX AAA11
XX AAA11
XX AAA11
XX AAA11
XX Ident

25-OCT-2000 AAA11232;

(first entry)

AAA11232 standard; DNA; 10

RAPD primer U-12 for detecting Leishmania species.

PCR primer; amplification; epidemiology; vaccine; ss.

Identification; Leishmania; isoenzyme; amplicon; promastigote; diagnosis;

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10

CCAGGAAGCC 1

1484 CCAAGAAGCC 1493

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Matches
                   Query Match
Best Local
                                                                                                                                                                                                                                                       AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of
                                                                                                                     e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeuti agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for adoptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
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                                                                      Sequence 10 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 90; 219pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated polynucleotides differentially expressed between metastatic non-metastatic breast cancer cells, useful for diagnosis, prevention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Roberts BL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            treatment of cancer.
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(ROBE/)
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) ROBERTS B L
) SHANKARA S.
l Similarity
9; Conserv
     Conservative
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98US-0090039P.
98US-0090040P.
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                                                                      A; 3 C; 4
                  28.0%;
90.0%;
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   0; Mismatches
                                                                      G; 3 T; 0 U; 0 Other;
                       Pred.
                                     Score
              2 8 . 4 ;
56 ;
                                       DB 1;
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     Indels
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RESULT 90
AAA633
XX AAA633
XX AAA633
XX AAA633
XX AAA633
XX AAA633
XX Human;
XX Human;
XX Homo 8
XX Cancer
XX Homo 8
XX Cancer
XX Y
PD 31-MAY
XX 21-NOV
PR 24-NOV
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PX VAX Velcul
XX Velcul
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   identifying isoenzymes and/or amplicons from, respectively, proteins or DNA extracted from cultured promastigotes derived from an isolate. The method uses a limited set of isoenzymes and/or amplification primers that are discriminatory for characteristics of a broad spectrum of Leishmania species. The isoenzymes used for detection include Glucose Phosphate Isomerase (GPI), Mannose Phosphate Isomerase (MPI), Nucleoside Hydrolase substrate deoxyinosine (NHD) and Phosphoglucomutase (PGM). The amplification primers used for PCR identification include primers AAA11232. The method is used to identify specific Leishmania species, either for diagnosis or for selection of appropriate treatments, also in epidemiological studies and for preparation of pharmaceuticals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying species of Leishmania, useful e.g. selection of treatment, using a limited set of amplification primers for differentiation.
              WPI; 2001-367706/38
                                                                                                  24-NOV-1999;
                                                                                                                              21-NOV-2000; 2000WO-US031922
                                                                                                                                                          31-MAY-2001.
                                                                                                                                                                                                                                           Human; transcriptome; gene expression pattern; cancer; drug cancer diagnosis; cell specific gene expression; ss.
                                                                                                                                                                                                                                                                                    Human colon epithelium specific transcriptome sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                  20-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                         AAH63301 standard;
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                                           Velculescu VE,
                                                                                                                                                                                       WO200138577-A2
                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a method for identifying Leishmania species
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                                                                       UNIV JOHNS HOPKINS
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                                         Vogelstein
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Pred. No.
                                           Kinzler
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56;
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RESULT 91
AAH63452
ID AAH63
XX Human
XX Human
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AAH63452
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XX AAH634
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                                                                                                                                          The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
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Sequence 10
                                                       cancer specific gene expression, standardise expression and restore the function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                               New isolated polynucleotides, useful for identifying specific cell type, such as cancer cell, comprises transcriptomes expressed in particular
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diagnosis; cel:
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   BP; 4 A; 4 C; 1
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cell specific gene expression; ss.
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90.0%;
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Pred. No. 56;
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Query Match

28.0%;

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RESULT 93
ABA06059/c
ID ABA060
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AC ABA060
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DT 10-JAN
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                                                                                                                                                                                                                                                                        cancer specific gene expression, standardise expression and restore the function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention
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              Human normal
                                      10-JAN-2002
                                                                                      ABA06059 standard; cDNA; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-NOV-1999;
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                                                                                                                                                                                                               Local
                                                                                                                                                                           1490 AGCCAGACTT 1499
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9; Conserv
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                                                                                                                                                                                                    Conservative
           hepatocyte expression gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-00448480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  expressed transcriptome sequence SEQ ID NO:
                                                                                                                                                                                                              28.0%;
90.0%;
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                                                                                      ВP
                                                                                                                                                                                                  0
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                                                                                                                                                                                                                         Score 8.4;
                                                                                                                                                                                                               Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kinzler
                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                               No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ₩,
            CDNA,
                                                                                                                                                                                                                         DB 1; Length 10;
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            SEQ ID NO: 36
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ARESULT 94
AAF38816
ID AAF38
AC AAF38
XX Yeast
XX Yeast
KW Yeast
KW nor 1
KW serik
KW nor 1
KW Serik
KW 11nk
XX Sacch
XX Sacch
YN WO200
XX 1-Di
XX 1-Ji
XX 14-Ji
XX 14-Ji
XX Yeast
YH Yelci
XX Yeast
YH Yeast
XX Yeast
YH Yeast
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a human normal hepatocyte expression gene group comprising 200 genes in the human normal hepatocyte. The cDNA of each gene comprises one of 200 fully defined nucleotide sequences as given in the specification. The gene group and the cDNAs corresponding to each of the genes in the group are useful in the diagnosis and treatment of human hepatopathy. The present sequence is a cDNA corresponding to a gene expressed by normal human hepatocytes
                                                                                                                                                                                                                                                 Saccharomyces cerevisiae.
                                                                                                                                                                                                                                                                                        Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
                                                                                                                                                                                                                                                                                                                                               Yeast NORF gene SAGE tag oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                          23-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                               AAF38816 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 10 BP; 1 A; 3 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 6; 26pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-JAN-2000; 2000JF-00023170.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-JAN-2000; 2000JP-00023170.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; hepatocyte;
                                                                                Velculescu V,
                                                                                                                                      16-JUN-1999;
                                                                                                                                                                14-JUN-2000; 2000WO-US016223
                                                                                                                                                                                            21-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                      WO200077214-A2
                                                                                                                                                                                                                                                                             linker; PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity nes 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1489 AAGCCAGACT 1498
                                                      2001-061874/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001-629566/73
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                                                                                                            UNIV JOHNS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGGCCAGACT
                                                                                                                                                                                                                                                                             primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                Vogelstein B,
                                                                                                                                      99US-00335032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene expression; hepatopathy;
                                                                                                            HOPKINS
                                                                                                                                                                                                                                                                             ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred.
                                                                                 Kinzler K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
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                                                                                                                                                                                                                                                                                                                                                NO:5555
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Yeast gene coding sequences comprising NORF genes with serial gene expression (SAGE) tags, useful for studying, monitoring affecting phases of the cell cycle.

analysis of

Example; Page 198; 419pp; English.

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AAF39726
ID AAF39
XX AAF39
XX AAF39
XX AAF39
XX Yeast
XX Yeast
XX Yeast
XX Inke
XX Ink
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nor previously assigned open reading serial analysis of gene expression; a linker; PCR primer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; nor previously assigned expression; antifungal; tag; identification;
                                          WPI; 2001-061874/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200077214-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:6465
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                Velculescu V,
                                                                                                                                                                                                                                                                16-JUN-1999;
                                                                                                                                                                                                                                                                                                                                      14-JUN-2000; 2000WO-US016223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1486 AAGAAGCCAG 1495
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                                                                                                                                                                                      UNIV JOHNS HOPKINS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10 BP; 6 A; 1 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                            Vogelstein B,
                                                                                                                                                                                                                                                                9908-00335032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28.0%;
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Pred. No. 5
                                                                                                                Kinzler K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1;
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Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle. 230; 419pp; English.

The present invention describes an isolated DNA molecule comprising a CC coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORP) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also Cd described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log thase, S phase and G2/M; (2) a method (M2) for screening candidate CC antifungal drugs comprising: (a) contacting a test substance with a yeast Ccll; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human genes which are involved in cell cycle progression CC contiguous mucleotides of a NORF gene whose expression varies as in M1; cc contiguous mucleotides of a NORF gene whose expression varies as in M1; cc candidate drug shaving a characteristic effect on gene expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug and contacting a yeast cell of at least 1 NORF gene whose capressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs. AAF33268 to AAF33267 represent linkers and PCR primers used in the SAGE captage and for identification of the present invention.

AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.

Sequence 10 BP; 4 A; 2 C; 4 G; 0 T; 0 U; 0 Other;

Ś Best Local Similarity Matches 9; Conserv Query Match 1487 AGAAGCCAGA 1496 Conservative 28.0%; 90.0%; 0; Mismatches Score 8.4; I Pred. No. 56; DB 1; Length 10; 0, Gaps 0

RESULT 96 AAF40474

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AGGAGCCAGA 10

AAF40474 standard; DNA; 10 BP

AAF40474;

23-MAR-2001 (first entry)

Yeast NORF gene SAGE tag oligonucleotide SEQ IJ NO:7213

Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification

PCR primer;

WO200077214-A2

Saccharomyces

cerevisiae

21-DEC-2000.

14-JUN-2000; 2000WO-US016223.

16-JUN-1999; 99US-00335032

(UYJO) UNIV JOHNS HOPKINS

Velculescu V, Vogelstein B, Kinzler

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RESULT 97
AAF40855
ID AAF4(
XX AAF4(
XX Yeas
DE Yeas
XX Yeas
XX Pas
HW OOT
XW IIII
XX HOOT
XW OOT
XX IIII
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes an isolated DNA molecule comprising coding sequence of a yeast gene selected from a group of 745 NORF (1 previously assigned open reading frame; or nonannotated ORF) genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example; Page 257; 419pp; English.
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                                                                                                                                                                                                                                                                                                     Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                   nor previously assigned open reading frame; nonannotated ORF; serial analysis of gene expression; antifungal; tag; identifi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF40855;
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                                      16-JUN-1999;
                                                                                                       14-JUN-2000; 2000WO-US016223.
                                                                                                                                                                         21-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yeast NORF
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                                                                                                                                                                                                                                                                                                                                                                    PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene SAGE tag oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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Pred. No. 56;
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                                                                                                                                                                                                                                                                                                                                                                                                          tag; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NO:7594.
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RESULT 98 AAF35202

AAF35202 standard; DNA; 10

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Matches Query Match

1492 CCAGACTTCA 1501 1 CCAGAGTTCA

10

Local

Similarity 9; Conserv

Conservative

<u>0</u>

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Indels

0

Gaps

0

28.0%;

Score 8.4; D Pred. No. 56;

DB 1;

Length 10;

Sequence 10 BP; 3 A; 3 C; 2 G; 2 T; 0 U; 0 Other;

serial analysis of gene expression; linker; PCR primer; ds.

Saccharomyces cerevisiae

14-JUN-2000; 2000WO-US016223

21-DEC-2000 WO200077214-A2 Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE;

tag;

nor previously assigned open reading frame; nonannotated

Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:1941

23-MAR-2001

(first entry)

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CC The present invention describes an isolated DNA molecule comprising a CC coding sequence of a yeast gene selected from a group of 745 NORF (not CC previously assigned open reading frame, or nonannotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC comprising a SAGE (serial analysis of gene expression) tag. Also CC comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log CC phase, S phase and G2/M; (2) a method (M2) for screening candidate CC cell; and (b) monitoring expression of a NORF gene whose expression of CC varies as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for CC contiguous nucleotides of a NORF gene whose expression of CC contiguous nucleotides of a NORF gene which comprises at least 10 CC contiguous nucleotides of a NORF gene whose expression varies as in M1; CC contiguous nucleotides of a NORF gene whose expression in a CC yeast cell comprising contacting a candidate drug as a member of a CC peast cell comprising contacting a yeast cell with a candidate drug and modifier of a NORF gene whose expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug and cCC expressed genes may be used to study, monitor and affect phases of the cell cycle. The ccc years of the cell cycle and the comprise of the cell cycle. The coll cycle and for identify candidate drugs which affect the cell cycle. The coll cycle and for identify candidate drugs which affect the cell cycle. The coll cycle and for identify candidate drugs which affect the cell cycle. The coll cycle and for identify candidate drugs which affect the cell cycle. The cycle and for identify candidate drugs which affect the cell cycle. The cycle and for identify candidate drugs which affect the cell cycle and the cycle and cycle. The cycle and cycle and cycle and cycle a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                coding sequence of a yeast gene selected from a group of 745 NORF (not CC previously assigned open reading frame; or nonannotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log CC phase, S phase and G2/M; (2) a method (M2) for screening candidate CC ell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for CC edl; and (f) monitoring expression of a NORF gene whose expression of CC varies as in: M1, where a test substance which modifies the expression of CC contiguous nucleotides of a NORF gene whose expression of CC contiguous nucleotides of a NORF gene whose expression of CC contiguous nucleotides of a NORF gene whose expression in a CC class of drugs having a characteristic effect on gene expression in a CC class of drugs having a characteristic effect on gene expression in a CC class of drugs having a characteristic effect on gene expression in a CC class of of custody and affect phases of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The CC methods may be used as markers of phases of the present invention. CC characteristic carries of an CC part of identification of antifungal drugs. ARA33268 to AAR44064 crepresent SAGE tags used in the exemplification of the present invention. CC characterism of the cream of the cell cycle. The cream of the characterism of the present invention. CC characterism of the cream of the coll of the SAGE content of the carries of the cell cycle. The cream of the characterism of the present invention.
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Best Local S
Matches 9
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AAF36577
                                                                                                   nor previously assigned open reading frame; nonamnotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
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                                                            Saccharomyces cerevisiae
                                                                                                                                                                    Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;
                                                                                                                                                                                                             Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:3316
                                                                                                                                                                                                                                                          23-MAR-2001
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                                                                                                                                                                                                                                                       (first entry)
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90.0%;
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RESULT 100
AAP36581/c
ID AAP365
XX AAP365
XX AAP365
XX C
DT 23-MAR
XX Yeast
XX Yeast;
XW Yeast;
XW nor pr
XW serial
KW serial
KW sechal

AAF36581;

23-MAR-2001

(first entry)

AAF36581 standard; DNA; 10

Yeast NORF gene SAGE tag oligonucleotide SEQ

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NO:3320

Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;

open reading frame; nonannotated

antifungal;

tag; identification;

expression;

Saccharomyces cerevisiae

nor previously assigned serial analysis of gene

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10

CTTGAGCAGC 1

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1497 CTTCAGCAGC 1506

Query Match Best Local S Matches 9

Similarity 9; Conserv

Conservative

<u>,</u>

Mismatches

1:

0

Gaps

0

28.0%; 90.0%; 3 C; 3

Score 8.4; Pred. No. 5

DB 1; Length 10, Other;

Sequence

10

BP; 2 A;

G; 2 T; 0 U; 0

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The present invention describes an isolated DNA molecule comprising a CC coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC certification of gene expression) tag. Also CC comprising administering a NORF gene whose expression varies by at CC cycle comprising administering a NORF gene whose expression varies by at CC cycle comprising and true the cell cycle selected from log thase. S phase and G2/M; (2) a method (M2) for screening candidate CC cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for cell cycle progression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human plnA with a probe which comprises at least 10 comprising contacting human DNA with a probe which comprises at least 10 comprising contacting a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a CC cass of drugs having a characteristic effect on gene expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug and monitor and affect phases of the call cycle, the differentially capter of the cell comprise of a comprise of the cell cycle. The comprise of the cell cycle and for identify candidate drugs which affect the cell cycle and for identify candidate drugs which affect the cell cycle and for identify candidate drugs which affect the cell cycle and the comprise used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
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of.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate CC ell; and (b) monitoring expression of a NORF gene whose expression of cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human penes which are involved in cell cycle progression of comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a cyeast cell comprision in the yeast cell of at least 1 NORF gene whose expression is study, monitor and affect phases of the cell cycle, the differentially corporated may be used to identify candidate drugs which affect the cell cycle and for identification of the cell cycle. The methods may be used to identify candidate drugs. AAF33268 to AAF34064 crepresent SAGE tags used in the exemplification of the cell cycle. The cycle and for identification of the the present invention. AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
                                                                                                                                                                                                                                                          RESULT 101
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Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes an isolated DNA molecule comprising coding sequence of a yeast gene selected from a group of 745 NORF (repreviously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also
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                                                                                   Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:5639
                                                                                                                               23-MAR-2001
                                                                                                                                                                         AAF38900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 10
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                                                                                                                                                                                                                AAF38900 standard; DNA; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                          Match 28.0%;
Local Similarity 90.0%;
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                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 8.4; DB 1; Length 10
                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. 56;
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                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
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RESULT 102
AAF33314
ID AAF333
XX
AC AAF333
XX
DT 23-MAR
XX
DE Yeast
XX

AAF33314 standard; ENA; 10

ВP

23-MAR-2001 (first entry)

Yeast putative coding sequence SAGE tag oligonucleotide SEQ ID NO:53

Query Match
Best Local Similarity
"---hes 9; Conserv

Conservative

0;

Mismatches

Indels

0

Gaps

0

90.0%;

Score 8.4; D. Pred. No. 56;

DB 1; Length 10; <u>,</u>

1486 AAGAAGCCAG 1495

Sequence 10 BP; 4 A; 2 C; 4 G; 0 T; 0 U; 0 Other;

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AAGGAGCCAG 10

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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not comprising a SAGE (serial analysis of gene expression) tag. Also comprising a SAGE (serial analysis of gene expression) tag. Also corporated are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at cell east 10% between any two phases of the cell cycle selected from log cantifungal drugs comprising: (a) a method (M2) for screening candidate cell; and (b) monitoring expression of a NORF gene whose expression of cell; and (b) monitoring expression of a NORF gene whose expression of cell; and (b) muman genes which are involved in cell; cycle progression cell; and (b) muman genes which are involved in cell; cycle progression cell; and (b) muman genes which are involved in cell; cycle progression centiquous mucleotides of a NORF gene whose expression varies as in M1; centifying human genes which are involved in cell; cycle progression in cell; cycle progression in a class of drugs having a characteristic effect on gene expression in a cyeast cell of at least 1 NORF gene whose expression in a cyeast cell of at least 1 NORF gene may be used to study, monitor and affect phases of the cell cycle. The centially cycle and for identify candidate drugs which affect the cell cycle. The methods may be used as markers of phases of the cell cycle. The centially cycle and for identify candidate drugs which affect the cell cycle and cycle and for identify candidate drugs. AAF33268 to AAF44064 ceptication of the present invention.

CC and (4) a method (4) a method (4) a method of antifungal drugs. AAF33268 to AAF44064 ceptication of the cycle and for identify candidate drugs. AAF33268 to AAF44064 ceptication of the present invention.
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of.

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RESULT 103
AAF34250
ID AAF342
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AC AAF342
XX
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AC AAF342
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                                                                                                                                                                                                                                                                                                                                         The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonamnotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also comprising a SAGE (serial analysis of gene expression) tag. Also correct the cell cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log thase, S phase and G2/M; (2) a method (M2) for screening candidate cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human DNA with a probe which comprises at least 10 comprising contacting human DNA with a probe which comprises as in M1; contiguous nucleotides of a NORF gene whose expression in a CC candidate antifying a candidate drug as a member of a CC class of drugs having a characteristic effect on gene expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially cycle and for identification of the cell cycle. The cell cycle and for identification of the cell cycle. The cell cycle and for identification of the cell cycle. The cell cycle and for identification of the cell cycle. The cell cycle and for identification of the present invention.

AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE comprises the exemplification of the present invention.
                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
  23-MAR-2001
                                                                                                                                                                                                                                                                                                                                       Sequence 10 BP; 4 A; 3 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Velculescu V,
                                         AAF34250;
                                                                              AAF34250 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-061874/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYJO ) UNIV JOHNS HOPKINS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linker; PCR primer;
                                                                                                                                                                                                                     1480
                                                                                                                                                                               μ.
                                                                                                                                                                                                                     ACGACCAAGA 1489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 22; 419pp;
                                                                                                                                                                                                                                                            Conservative
  (first entry)
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                                                                                                                                                                                                                                                                             28.0%;
90.0%;
                                                                              BP.
                                                                                                                                                                                                                                                        ٥,
                                                                                                                                                                                                                                                                             Score 8.4; D
Pred. No. 56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kinzler K;
                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                               DB 1; Length 10
                                                                                                                                                                                                                                                        1;
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                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                        0;
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RESULT 104 AAF35602/c ID AAF356 XX

AAF35602 standard; DNA; 10

밁 Ś

Matches

Conservative

0 Pred. No. Score 8.4;

Mismatches

1:

0

Gaps

0

DB 1;

Length 10; Indels

28.0%; 90.0%;

1480 ACGACCAAGA 1489 Similarity 9; Conserv Sequence 10

BP; 4 A; 3 C; 3 G; 0 T; 0 U; 0 Other;

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CC coding sequence of a yeast gene selected from a group of 745 NORF (not Cpreviously assigned open reading frame; or nonannotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC described are; (1) a method (M1) of using NORF genes to affect the cell Cycle comprising administering a NORF gene expression tag. Also CC described are; (1) a method (M1) of using NORF genes to affect the cell Cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate cc antifungal drugs comprising; (a) contacting a test substance with a yeast CC varies as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human DNA with a probe which comprises at least 10 cc comprising contacting human DNA with a probe which comprises at least 10 cc contiguous nucleotides of a NORF gene whose expression varies as in M1; cc cand (4) a method (M4) for identifying a candidate drug as a member of a cyeast cell comprising contacting a yeast cell with a candidate drug and and comprising contacting a yeast cell with a candidate drug and and comprising contacting a yeast cell of at least 1 NORF gene whose expression in a cyeast cell of at least 1 NORF gene may be used to study, monitor and affect phases of the cell cycle. The central cycle and for identification of the cell cycle. The correspond to study, monitor and affect phases of the cell cycle. The correspond to the cell cycle where the cell cycle and for identification of the central cycle. The central cycle and for identification of the central cycle and correspond to the cell cycle. The central cycle and for identification of the central cycle and correspond to the cell cycle. The cycle and cycle a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-061874/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example; Page 35; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Velculescu V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9908-00335032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kinzler K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ID NO:989
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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene expression) tag. Also comprising a SAGE (serial analysis of gene expression) tag. Also coycle comprising administering a NORF gene whose expression varies by at coycle comprising administering a NORF gene whose expression varies by at coycle comprising administering a NORF gene whose expression tog contain and the probabes of the cell cycle selected from log contains and G2/M; (2) a method (M2) for screening candidate cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for contiguous nucleotides of a NORF gene which modifies the expression of comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression of and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression in a green expression in a specific to gene expression in a green expression in a description of the cell cycle, the differentially cycle and for identification of antifungal drugs. AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention.

CC methods may be used as markers of phases of the present invention. AAF33267 represent linkers and PCR primers used in the SAGE method, in the PCR present linkers and PCR primers used in the SAGE method. In the PCR present linkers and PCR primers weed in the SAGE method.
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                                                                             Query Match
Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Saccharomyces cerevisiae
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                                                                                                                                                             Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Page 83; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-061874/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-JUN-1999;
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  10
                                                                               Similarity 9; Conser
                                                                                                                                                                                                     in the exemplification of the present invention
                                       CTTCAGCAGC 1506
                                                                                                                                                               BP;
                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                             2 A;
                                                                                                                                                             3 C;
                                                                                                  28.0%;
                                                                                                                                                             3 G;
                                                                                                  Score 8.4;
Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kinzler
                                                                                                                                                             2 T; 0 U;
                                                                               Mismatches
                                                                                                                     DB 1; Length 10;
                                                                                                                                                               0 Other;
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RESULT 105

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> 1486 AAGAAGCCAG 1495 l Similarity 9; Conserv

Conservative

28.0%; 2 C; 4

Score 8.4; D. Pred. No. 56;

DB 1; Length 10

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0

Gaps

0

AAGAAGCCGG

10

Query Match Best Local

Sequence

10

BP;

4. A;

G; 0 T; 0 U; 0 Other;

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Coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not comprising a ssigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag, Also coycle comprising administering a NORF gene whose expression varies by at Cleast 10% between any two phases of the cell cycle selected from log containing administering a NORF gene whose expression varies by at Cleast 10% between any two phases of the cell cycle selected from log cantifungal drugs comprising; (a) a method (M2) for screening candidate cell; and (b) monitoring expression of a NORF gene whose expression of cell; and (b) monitoring expression of a NORF gene whose expression of cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for centiguous nucleotides of a NORF gene whose expression of comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell of at least 1 NORF gene whose expression in the yeast cell of at least 1 NORF gene whose expression in the past cell of at least 1 NORF gene may be used to study, monitor and affect phases of the cell cycle. The methods may be used as markers of phases of the cell cycle. The cycle and for identification of antifungal drugs. AAF33268 to AAF33267 represent 1 inkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kinzler K;
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CC comprising a SAGE (a) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast CC antifungal drugs comprising: (a) contacting a test substance with a yeast CC varies as in M1, where a test substance which modifies the expression CC the yeast gene is a candidate antifungal drug; (3) a method (M3) for CC contiguous nucleotides of a NORF gene whose expression CC contiguous nucleotides of a NORF gene whose expression varies as in M1; (2) a method (M4) for identifying a candidate drug as a member of a CC class of drugs having a characteristic effect on gene expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug and CC monitoring expression in the yeast cell of at least 1 NORF gene whose expression is study, monitor and affect phases of the cell cycle, the differentially CC expressed genes may be used as markers of phases of the cell cycle. The cycle and for identification of the cycle, the differentially cycle and for identification of the present invention.

CC AAF33262 to AAF3367 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 106
AAF41495/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Saccharomyces cerevisiae
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; Page 294; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         affecting phases of the cell cycle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-061874/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Velculescu V;
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Query Match Best Local S

10 ВP;

0 ?

N

G; 4 T; 0 U; 0 Other;

Similarity 9; Conserv

Conservative

0

Mismatches 56 DB 1; ۲.

0

Gaps

0

Query Match Sequence 10

BP; 5 A; 2 C; 2 G; 1 T; 0 U;

0 Other;

28.0%;

Score 8.4;

DB 1; Length 10;

28.0%; 4 C;

Score 8.4; Pred. No. 5

Length 10; Indels

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The present invention describes an isolated DNA molecule comprising a CC coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genees CC comprising a SAGE (serial analysis of gene expression) tag. Also CC comprising a SAGE (serial analysis of gene expression) tag. Also CC cycle comprising administering a NORF genee whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log CC phase, S phase and G2/M; (2) a method (M2) for screening candidate CC cell; and (b) monitoring expression of a NORF gene whose expression of CC varies as in M1, where a test substance which modifies the expression CC varies as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for contiguous nucleotides of a NORF gene whose expression of CC contiguous nucleotides of a NORF gene whose expression varies as in M1; and (b) munan genes which are involved in cell cycle progression in a CC contiguous nucleotides of a NORF gene whose expression varies as in M1; and (d) a method (M4) for identifying a candidate drug as a member of a CC class of drugs having a characteristic effect on gene expression in a CC vertice and (d) for identifying a yeast cell with a candidate drug and cc centroly, monitor and affect phases of trugs. The NORF gene may be used to study, monitor and affect phases of the cell cycle. The cc centroly may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF33268 to AAF34064 CC represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:1165
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RESULT 108
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                                                       CC coding sequence of a yeast gene selected from a group of 745 NORF (not coronary sequence of a yeast gene selected from a group of 775 NORF (not previously assigned open reading frame; or nonannotated ORP) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC described are: (1) a method (M1) of using NORF genes to affect the cell Cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate CC antifungal drugs comprising: (a) contacting a test substance with a yeast Cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for CC identifying human genes which are involved in cell cycle progression of the yeast gene whose expression varies as in M1; contacting human DNA with a probe which comprises at least 10 CC contiguous nucleotides of a NORF gene whose expression varies as in M1; characteristic celfect on gene expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug and CC monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially cyrosesed may be used as markers of phases of the cell cycle. The cell cycle and for identification of antifungal drugs which affect the cell cycle, the differentially cyrosesent sAGE tags used in the exemplification of the present invention. CC method, in the exemplification of the present invention.
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                                    exemplification of the present invention
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                                                                                                                                                                                                                                                                 (SNPs) in the human growth hormone releasing hormone receptor (GHRHR) gene located on chromosome 7p14, and methods for haplotyping and/or genotyping the GHRHR gene. The methods of the invention make use of allele-specific oligonucleotides (ASOs) as probes and primers and/or primer-extension oligonucleotides for detecting the GHRHR gene polymorphisms. The polynucleotides and screened compounds are useful the treatment of diseases associated with GHRHR activity, such as isolated growth hormone deficiency (IGHD) and pituitary adenomas. AAS19648-AAS19673 represent primer-extension oligonucleotides for detecting human GHRHR gene polymorphisms
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                                                                                                                                                                                                                     Sequence 10 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genotyping human Growth hormone releasing hormone receptor gene of individual for determining haplotype of individual by determining identity of nucleotide pair at specific polymorphic sites for two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-APR-2000; 2000US-0197978P
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GAAGCCTGAC 10
                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention relates to novel single nucleotide polymorphisms
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                                                                                                                                   Score 8.4; D
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Pred. No. 5
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                                                                                                            Mismatches
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                                                                                                                                                              DB 1;
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                                                                                                                                                                 Length 10
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                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                            are useful for
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                                                                                                            Gaps
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ABL52200 standard;

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RESULT 111
ABK81399/c
ID ABK813
XX
AC ABK813
XX
DT 13-AUG
DE SCYA21
XX
XX
DX SMBA11
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                                                                                                                                                                                                                                                                                                                                                                The present invention describes an isolated human period (Drosophila) (C) homologue 1, (PER1) polynuclectide (I) comprising a sequence which is a copolymorphic variant for a reference sequence (ABL52077) for the PER1 gene (C) or its fragment, or a polymorphic variant of a reference sequence (ABL52078) for a PER1 cDNA or its fragment. The present invention also (C) describes methods for genotyping and haplotyping the PER1 gene of an individual (I) is useful in studying the expression and function of PER1, and in expressing PER1 protein for use in screening for candidate (C) drugs to treat diseases related to PER1 activity. (I) is useful for therapeutic purposes A recombinant non-human organism transformed or transfected with (I) can be used for studying expression of the PER1 (I) can be used for studying expression of the PER1 (I) can be used for studying and testing of drugs targeted against PER1 protein, and for testing the efficacy of therapeutic agents and compounds for disorders associated with circadian rhythm regulation. The present sequence represents a preferred oligonucleotide primer for them.
                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; period (Drosophila) homologue 1; PER1; polymorphic variant; polymorphic site; genotyping; haplotyping; circadian rhythm regulation; single nucleotide polymorphism; SNP; gene; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel isolated human period Drosophila homolog 1 polynucleotide, useful for therapeutic purposes, for studying the expression and function of the polynucleotide, and for expressing the homolog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
Small inducible cytokine subfamily A (Cys-Cys) member 21; SCYA21;
                              SCYA21 gene primer extension oligonucleotide #3
                                                                    13-AUG-2002
                                                                                                      ABK81399
                                                                                                                                      ABK81399 standard;
                                                                                                                                                                                                                                                                                                                                                              Sequence 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-MAR-2002
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                                                                                                                                                                                                                                                           1497 CTTCAGCAGC 1506
                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                           CGTCAGCAGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 16; 162pp; English.
                                                                                                                                                                                                                                                                                                                                                            BP; 2 A; 4 C; 3 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                  which
                                                                                                                                                                                                                                                                                              Conservative
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                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                  is used in the
                                                                                                                                      DNA;
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90.0%;
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                                                                                                                                                                                                                                                                                                                              Score 8.4;
                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                exemplification of the present invention
                                                                                                                                                                                                                                                                                                          No.
                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                            1.
                                                                                                                                                                                                                                                                                                                            Length 10;
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drug
                                                                                                                                                                                        New genetic variants comprising haplotypes of the small inducible cytokine subfamily A, member 21 (SCYA21) gene, useful in improving the efficiency of screening for drugs for treating immunological disorders
The invention describes an isolated polynucleotide, which comprises genes and haplotypes of the small inducible cytokine subfamily A (Cys-Cys),
                                                                                                  Claim 16; Page 13; 56pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-OCT-2001; 2001WO-US046141.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymorphism; haplotype; immunological disorder; gene expression;
drug development; immunomodulator; primer extension; oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                        WPI; 2002-435528/46
                                                                                                                                                                                                                                                                                                                                                                                    Bentivegna SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-OCT-2000; 2000US-0241622P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GENA-) GENAISSANCE PHARM INC
                                                                                                                                                          iciency of screening for drugs targeting SCYA21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sapiens
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member 21 (SCYA21) gene. The polynucleotide comprises polymorphic sites referred to as PS1-5 to designate the order in which they are located in the gene. The polymorphisms and haplotypes of SCYA21 gene are useful for validating whether SCYA21 is a suitable target for drugs to treat immunological disorders and disorders associated with its abnormal expression or function, screening for such drugs and reducing bias in clinical trials of such drugs. Haplotype information would be useful in improving the efficiency and output of several steps in the drug discovery and development process, including target validation, identifying lead compounds and early phase clinical trials. The methods are useful in screening for compounds targeting SCYA21 to treat a specific condition or disease predicted to be associated with SCYA21 activity, e.g. immunological disorders. This sequence represents a primer extension oligonucleotide used to identify polymorphic sites in the Sequence 10 BP; 0 A; 4 C; 2 G; 4 T; 0 U; 0 Other; gene В ۳. Length 10;

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ABK96063/c
ID ABK960
XX
AC ABK960
AC ABK960
DT 24-SEP
XX
DE Human
XX
Human

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Best Local
                                                                                                                                                                                                                                                                                                                                         Human; lipase; hormone sensitive; LIPE; isogene; obesity; male sterility;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK96063;
                                                                                                                                                                                                                                                                                           polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Human LIPE gene polymorphism detection oligonucleotide primer #38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK96063 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1487 AGAAGCCAGA 1496
                                                                                                                                                                                                sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                           primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
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23-MAY-2002 WO200240502-A2

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RESULT 113
AAD25204
ID AAD252
XX AAD252
AC AAD252
XX Human;
KW Human;
KW Haplot
KW Homo &
XX Hom
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a new polynucleotide comprising a nucleotide sequence which comprises lipase, hormone sensitive (LIFE) isogenes. The invention is useful in screening for drugs targeting LIFE isogenes that are useful for treating obesity and male sterility. The lipase is the invention are useful for improving the efficiency and reliability of several steps in the discovery and development of drugs for treating diseases associated with LIFE activity. The polynucleotide is useful in studying the expression and function of LIPE, and in expressing LIPE protein for use in screening for candidate drugs to treat diseases related to LIPE activity. It is also useful in studying the effect of the variation on the biological activity of LIPE as well as on the binding affinity of candidate drugs targeting LIPE for the treatment of obesity and male sterility. The invention is useful for studying the expression of LIPE isogenes in vivo, for in vivo screening and testing of thrust sargeted against LIPE protein, and for testing the efficacy of the treatment of activity and male sterility in a biological system. The present nucleic acid sequence represents one of a collection (ABK96026-ABK96083) of oligonucleotide primers that were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                     Human; homeo box D3; HOXD3; polymorphism; developmental disorder;
haplotype; HT; allele-specific oligonucleotide; ASO; tumour; therapy;
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                                                            25-MAY-2000; 2000US-0207076P.
                                                                                                                        24-MAY-2001; 2001WO-US016982.
                                                                                                                                                                                       29-NOV-2001
                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human homeo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAD25204;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAD25204 standard; DNA; 10
      (GENA-)
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                                                                                                                                                                                                                                                                                                                                                                     screening;
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      GENAISSANCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention to detect polymorphisms in the human LIPE gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          box D3 (HOXD3) gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                     cytostatic; primer;
      PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28.0%;
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Pred. No. 56;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polymorphism detecting primer
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The invention relates to genetic variants of the homeo box D3 (HOXD3) CC gene. HOXD3 gene includes 9 polymorphic sites PS1-PS9. Haplotypes (HTS) CC or haplotype pairs (HP) for PS1-PS9 in the HOXD3 gene are useful for improving the efficiency and reliability of several steps in the CC discovery and development of drugs for treating diseases associated with CC discovery and the expression and function of HOXD3 isogene is useful in studying the expression and function of HOXD3 and in CC expressing HOXD3 protein for use in screening for candidate drugs to treat diseases related to HOXD3 activity and in studying the effect of the variation on the biological activity and in studying the effect of CC binding affinity of candidate drugs targeting HOXD3 as well as on the CC binding affinity of candidate drugs targeting HOXD3 for the treatment of CC developmental disorders and tumours. An antibody against HOXD3 is useful in a variety of diagnostic and prognostic formats and therapeutic CC methods. A recombinant non-human organism is useful in studying CC expression of the HOXD3 is useful as probes and primers and for assaying a polymorphism in the target region. The present sequence is a primer candidate drugs target region. The present sequence is a primer candidate drugs target region.
Sequence 10 BP; 4 A; 3 C; 2 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 18; Page 13; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         developmental disorders and tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-075363/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the protein, and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  genetic variants of Homeo Box D3 for studying expression and function the protein, and for screening drugs to treat diseases e.g.
                                                   detecting human HOXD3 gene polymorphisms
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Matches Query Match Best Local 1499 1 TCAGCAGACA 10 Similarity 9; Conserv TCAGCAGCCA 1508 Conservative 28.0%; 0; Mismatches Score 8.4; D Pred. No. 56; DB 1; Length 10 1. Indels 0 Gaps

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RESULT 114
ARVS4823/C
ID 4BVS48
XX ABV848
XX ABV848
XX BAGE t
KW CH; li
KW CH; li
KW expres
XX JP2002
XX WPI; 2
DAN
XX WPI; 2
XX Claim ABV84823; ABV84823 standard; cDNA; 10 Human haemopexin 12-DEC-2002 (first entry) SAGE tag #633. ВP

SAGE tag; serial analysis of gene expression; human; chronic hepatitis CH; liver tissue; hepatocellular carcinoma; cancer; tumour; HCC; expression pattern; 88.

Homo sapiens.

30-JUL-2002 JP2002209591-A.

19-JAN-2001; 2001JP-00012328

19-JAN-2001; 2001JP-00012328

KAGAKU GIJUTSU SHINKO JIGYODAN

WPI; 2002-631294/68

Human chronic hepatitis C tissue expression exasperating gene group comprises 100 high-ranking genes.

Claim 55; Page 28; 139pp; Japanese

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RESULT 115
ABK54423/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; 88; primer; ISL1; islet-1; chromosome 5q; motor neuron defect; diabetes; transcription factor; LIM; homeodomain; antidiabetic; PCR; gene therapy; primer extension.
                                                                                                                                                          expressing ISL1 treat diabetes.
                                                                                                                                                                          Novel isolated polynucleotide which is a polymorphic variant of ISL1 transcription factor, LIM/Homeodomain, (islet-1) (ISL1) used for expressing ISL1 protein isoform and for screening drug candidates to
                                                                                                                                                                                                                                                                                                                                                                                                               04-AUG-2000; 2000US-0223535P
                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-AUG-2001; 2001WO-US024664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human ISL1 gene ASO primer extension sequence #18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK54423 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 10 BP; 1 A; 3 C; 3 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expressed in chronic hepatitis C liver tissue
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                                                                                                                                                                                                                                                                                                                                                                      GENAISSANCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAGCCAGACT 1498
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                                                                                                                                                                                                                                                                                                                    Koshy B,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                      PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
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Pred. No. !
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The invention relates to an isolated polynucleotide sequence which comprises ISI1 transcription factor (islet-1, of the LIM/Homeodomain family), isogene and the polymorphic variants of the coding region (cDNA). Also included are a recombinant non-human organism expressing

The present invention relates to a new polypeptide comprising a sequence which is a polymorphic variant of a reference sequence for ADRB3 (beta-3-adrenergic receptor) protein. The reference sequence comprises a sequence

14;

90pp;

English

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RESULT 116
ABK11491
ID ABK114
XX ABK1149
AC ABK114
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XX O1
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                                                                                                                                                                                      Novel genetic variants of beta-3-adrenergic receptor gene useful in studying expression and function of the protein, and for screening drugs to treat diseases e.g. obesity, non-insulin dependent diabetes mellitus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; beta-3-adrenergic; receptor; ADRB3; primer; anorectic; ss; antidiabettc; gene therapy; morbid obesity; insulin resistance; non-insulin-dependent diabetes mellitus; haplotyping; SNP; single nucleotide polymorphism.
                                                                                                                                Claim 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide primer #3, to detect human ADRB3 gene polymorphisms
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                                                                                                                                                                                                                                                                                                                         WPI; 2002-241571/29.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-JUN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GENA-) GENAISSANCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1495
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                                                                                                                            Page 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 2 A; 3 C; 3 G; 2 T; 0 U; 0 Other;
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                                                                                                                         91pp; English
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Pred. No. 56;
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RESULT 117
ABK72629
ID ABK7262
AC ABK726
AC GENA-
AC GENA-
AC ABK726
AC ABK7
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                                                     The invention describes an isolated human leukotriene B4 receptor (chemokine receptor-like 1) (LTB4R) polynucleotide (I) comprising a sequence which is a polymorphic variant for a reference sequence for LTB4R gene or its fragment, or a polymorphic variant of a reference sequence for a LTB4R cDNA or its fragment. LTB4R is a potent chemoattractant that is primarily involved in inflammation, immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of 408 amino acrus us grant the polymorphic variant are
                                                                                                                                                                                                                                                                                                                                                                                   Novel isolated human leukotriene B4 receptor polynucleotide, useful therapeutic purposes, for studying expression and function of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bieglecki KM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200230949-A2
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                                 responses and host
                                                                                                                                                                                                                                                                                            Claim 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENA-) GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1484 CCAAGAAGCC 1493
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCAGGAAGCC 10
                                                                                                                                                                                                                                                                                         Page
                                                                                                                                                                                                                                                                                                                                              purposes, to,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               84
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                                                                                                                                                                                                                                                                                      14; 69pp;
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      defense against in function of LTB4R,
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                                                                                                                                                                                                                                                                                                                                                           receptor.
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                                    (I) is useful
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                                                        The invention relates to detecting CC (colorectal cancer e.g. colorectal adenoma), comprising: (a) detecting macrophage inhibitory cytokine (MIC) or renal dipeptidase (RDP) in faeces or blood of a subject and comparing amount of MIC or RDP detected to that in normal subjects, where an elevated amount of MIC or RDP in the subject is an indicator of CC in subject; (b) isolating mRNA sample from faeces of a subject, detecting MIC or RDP mRNA in the mRNA sample, and comparing amount of MIC or RDP mRNA in the subject is an indicator of CC in subject or RDP mRNA in the subject is an indicator of CC in subject; (c) isolating epithelial cells from blood of a subject, isolating an mRNA sample from faeces of a subject, isolating an mRNA sample from faeces of a subject is an indicator of CC in subject; (c) isolating epithelial cells from blood of a subject, isolating an mRNA sample from faeces of a subject or epithelial cells, detecting MIC or RDP mRNA in the mRNA sample, and comparing the amount of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in normal subjects, where
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  use in screening for candidate drugs to treat diseases related to LTB4R activity, e.g. inflammatory disorders recombinant non-human animal is useful for studying expression of the LTB4R isogenes in vivo, for in vivo screening and testing of drugs targeted against LTB4R protein, and for testing the efficacy of therapeutic agents and compounds for diseases associated with LTB4R activity, e.g. inflammatory disorders, in a biological system. This sequence represents a primer extension oligonuclection used for detecting polymorphisms in the leukotriene B4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detecting colorectal cancer in a subject, involves detecting macrophage inhibitory cytokine or renal dipeptidase or their mRNA in feces or blood of the subject.
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30-MAY-2002; 2002US-0383805P.
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9; Conserv
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of CC in the subject; (d)

or RDP mRNA

in

the mRNA sample is

indicative

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XX Nucle
DT 20-FEE
XX Nucle
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PN U020/
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XX HAM
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Best Local Similarity
Matches 9; Conserv
                                         The invention comprises a method for the PCR amplification of nucleic acids. The method involves a set of primers, where two of the primers in solution and at least two other primers are attached to a solid support. The method of the invention can be used for the analysis of a nucleic acid or a mixture of nucleic acids, including: single-stranded DNA molecules, double-stranded DNA molecules, double-stranded DNA molecules. The
                                                                                                                                                                                                                                                                                                                                                                                                      Amplification of nucleotide sequences from polynucleotides by chain extension of oligonucleotide primers, comprises 2 oligonucleotides in solution, 2 attached to supports and both share complementary sequences.
                                                                                                                                                                                                                                                                                                                          Disclosure; Fig 17; 60pp; English
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RAPD; PCR; primer; random amplified polymorphic DNA.
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90.0%;
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Pred. No. 56;
        a random amplified
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        polymorphic DNA (RAPD)
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RESULT 121 AAX14955/c ID AAX149 XX AC AAX149

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Matches 9
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Best Local S
Matches 9
                                                                                                                                                                                                                                                            The invention comprises a method for the PCR amplification of nucleic acids. The method involves a set of primers, where two of the primers in solution and at least two other primers are attached to a solid support. The method of the invention can be used for the analysis of a nucleic acid or a mixture of nucleic acids, including; single-stranded DNA molecules, double-stranded DNA molecules, double-stranded DNA molecules and mRNA molecules. The present DNA sequence represents a random amplified polymorphic DNA (RAF PCR primer of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 10
                                                                                                                                                                                                            Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Amplification of nucleotide sequences from polynucleotides by chain extension of oligonucleotide primers, comprises 2 oligonucleotides in solution, 2 attached to supports and both share complementary sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleic acid PCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RNA analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid amplification; nucleic acid analysis; DNA analysis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABT14323 standard;
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nes 9; Conserv
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                                            AAGAAGCCAG
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                                                                                                                                                                                                               BP; 4 A; 2 C; 4 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fig 17; 60pp; English.
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                                                                                                     Conservative
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                                                                                                                              Score 8.4; D
Pred. No. 56;
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Pred. No. 5
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                                                                                                                                                       Length 10;
                                                                                                       Indels
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                                                                                                  Gaps
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                                                                                                                                                                          RESULT 122
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Matches 9
                                                                                                                                                                                                                                                                                                                                                                                               sequences to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is designed to form a triple-strand structure with part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detection of oncogenes and Hepatitis B virus. (Updated on 27-AUG-2003 to correct OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents a polynucleotide that is able to form triple helix with a double stranded sequence. Cytosine bases in the present can be replaced with 5-methylcytosine for increased triplex stability. The present sequence is used in the assay of the invention, where it can be part of the anchor DNA or reporter DNA sequence. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-AUG-2003
24-MAR-1999
              Human; skin ageing; skin stress; EST; expressed sequence tag; ss
                                            Human skin stress/ageing related EST SEQ ID NO 309.
                                                                             10-SEP-2002
                                                                                                             ABQ86554;
                                                                                                                                            ABQ86554 standard; cDNA; 11 BP
                                                                                                                                                                                                                                                                                                                                                     Sequence 11
                                                                                                                                                                                                                                                                                                                                                                                     field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             assay comprises adding a sample containing double-stranded DNA test
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Col 25-26; 168pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Triplex formation; DNA detection; triple helix; identification; bacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Triple helix third strand of 23S rRNA gene nucleotides 486-496
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                                                                                                                                                                                                                                                                                      Local Similarity
les 9; Conserv
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                                                                                                                                                                                                                                                                                      Conservative
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                                                                            (first entry)
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(first entry)
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RESULT 123
ABQ86306/c
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AC ABQ863
XX IO-SEP
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         optionally translated, genetically encoded factors (A) obtained from young and aged skin, to identify that genes that show strong differential expression. (A) comprises protein or mRNAs or their fragments (M1) is useful for: identifying markers of skin ageing and/or stress; determining skin ageing and/or stress; and identifying or determining the effects of pharmaceutical or cosmettic agents for control of skin ageing. The present sequence is one of a group of human skin ageing/stress related expressed sequence tags (ABQ86246-ABQ87680) of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to identifying (M1) genes in vitro that, or animals, are important for skin ageing and/or skin stress by analysis of gene expression between mixtures of transcribed and
                                                                                                                                                                                                        03-JAN-2001; 2001DE-01000121
                                                                                                                                                                                                                                                                                                                                                                                      WO200253773-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; skin ageing; skin stress; EST; expressed sequence tag;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 8; Page 49; 325pp; German
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                Claim 8; Page 7/; JULY, JULY, JULY, JULY, In humans or animals, are important for skin ageing and/or skin stress by serial analysis of gene expression between mixtures of transcribed and optionally translated, genetically encoded factors (A) obtained from young and aged skin, to identify that genes that show strong differential expression. (A) comprises protein or mRNAs or their fragments. (M1) is useful for: identifying markers of skin ageing and/or stress; determining the effects of the present of the present of the present of skin ageing and/or stress; and identifying or determining the effects of the present of skin ageing and/or stress; and identifying or determining the present of the present of skin ageing. The present
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                                                                                                                                                                                                                                                      The invention relates to in vitro identification (MI) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (MI) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; sclerodermatichthyosis; atopic dermatitis; acne, sebornhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
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Pred. No. 60;
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Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                      disorders, specifically neurodermatitis; sumburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacca; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SACE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                  Human skin
                                                                                                  21-OCT-2002
                                                                                                                                ABV69450;
                                                                                                                                                                ABV69450 standard;
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Pred. No. 60
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RESULT 129
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                                                                                                                                                                                                                                                                                                                                                                                                                   Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human skin
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                                                               Disclosure; Page 139; 1345pp; German
                                                                                                            In vitro identification of a homeostasis and identifying
                                                                                                                                                              WPI; 2002-590638/63.
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RESULT 133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against e.g. skin cancer.
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                                          Homo sapiens
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RESULT 134
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Human;
KW Human;
KW immuno
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Homo 8
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vitro identification of skin-expressed genes, useful for determining
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homeostasis and identifying cosmetic or pharmaceutical agents against
    196;
1345pp;
    German
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The invention relates to in vitro identification (MI) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (MI) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin Sequence 11 disorders, specifically neurodermatitis; sunburn; psoriasis; sclerod ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of skin. The present sequence is that of a human expressed sequence tag (EST) of the invention BP; 3 A; 4 C; w G; 1 T; 0 U; 0 Other; scleroderma; tag of the

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Matches
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Best Local
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              CCAAGAAGCC
                            Conservative
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RESULT 135
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XX ABV686
XX ABV686
XX Human
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The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin

specifically

neurodermatitis;

sunburn;

scleroderma;

Disclosure; Page 203;

1345pp; German

In vitro identification of a homeostasis and identifying

of skin-expressed genes, useful ying cosmetic or pharmaceutical a

agents

for determining agents against

2002-590638/63.

D,

Conradt

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Hofmann

×

.g. skin

cancer.

03-JAN-2001; 2001DE-01000127 20-DEC-2001; 2001WO-EP015179

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RESULT 136
ABV63846/c
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                                                                                                                                                                                                                                                                            Disclosure; Page 69; 1345pp; German.
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                                                                             G; 2 T; 0 U;
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Pred. No.
                                      Score 8.4; D
Pred. No. 60;
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                          Mismatches
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RESULT 138
ABV66909/c
ID ABV669
XX
AC ABV669
AC ABV669
XX
DT 21-OCT
XX
DE Human
XX
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ARVEST19
ID ABV652
XX ABV652
XX ABV652
XX T1-OCT
XX Human
XX Human
XX Human
XX Human
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XX Homo s
XX WO2002
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disorders, specifically neurodermatitis; sunburn; psoriasis; sclerodermatichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma, basal cell carcinoma, and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; skin; dermatological; vulnerary; antipsoriatic; antis
immunosuppressive; antiinflammatory; cytostatic; SAGE; neuro
psoriasis; dermatitis; skin cancer; EST; expressed sequence
                               Human skin EST 4695
                                                                                    21-OCT-2002
                                                                                                                                          ABV66909;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (M1) is useful for identifying genes involved in skin homeostasis; determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 108; 1345pp;
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                                                                                                                                                                                               ABV66909
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HENK ) HENKEL KGAA
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                                                                                                                                                                                               standard;
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neurodermatitis;
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RESULT 139
ABV71267/c
ID ABV712
XX ABV712
XX 21-OCT
XX 21-OCT
XX Human;
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KW immuno
KW psoria
XX Homo s
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Best Local :
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                                                                                                                                                                  Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis;
 03-JAN-2001; 2001DE-01000127
                                20-DEC-2001; 2001WO-EP015179
                                                                                           WO200253774-A2
                                                                                                                        Homo sapiens.
                                                                                                                                                      immunosuppressive; psoriasis; dermati
                                                                                                                                                                                                                   Human skin EST 9053
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                                                              11-JUL-2002.
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                                                                                                                                                       dermatitis; skin cancer;
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Pred. No. 60;
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                                                                                                                                                       EST;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ed genes, useful for determini pharmaceutical agents against
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RESULT 140
ABV67181
ID ABV671
XX ABV671
XX ABV671
XX Human
XX Human
XX Human
XX Homo s
SY Homo s
YN WO2002
XX IN VILL
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Best Local S
Matches
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                                                e.g. skin cancer.
                                                                                                                                                                                                             03-JAN-2001; 2001DE-01000127
                                                                                                                                                                                                                                                                                                                                                                       immunosuppressive; antiinflammatory; cytostatic; SAGE; neuroderma psoriasis; dermatitis; skin cancer; EST; expressed sequence tag;
                                                               In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                         20-DEC-2001; 2001WO-EP015179
                                                                                                                                                                                                                                                                            11-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                          Human; skin;
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                                                                                                                2002-590638/63
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9; Conserv
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Disclosure; Page 162;

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                                                                                                                                                                                                                   The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin.
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                                                                                                       disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            e.g. skin
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The invention describes a method of directionally assembling a recombinant viral genome comprising ligating the subclones of the viral genome to assemble a recombinant viral genome, particularly coronavirus. For directionally assembling a recombinant viral genome. In particular, the method is useful for manipulating the genomes of higher plants and animals, as well as bacteria and viruses. In particular, the method is useful for the precise genetic manipulation of individual chromosomes in whole plants and animals and the construction of artificial chromosomes for gene therapy. The genomes produced are useful in preparing vaccines and expression vectors (e.g., TGE vectors and vaccines), which are useful in protocols involving vaccination, gene transfer and gene therapy. This sequence represents the interconnecting junction site C/DE-1 used in the

Example 7; Page 22; 42pp; English.

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RESULT 143
ABT16435/c
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                                                  The invention relates to a nucleic acid segment from the human neurokinin CC 1 receptor (TACR1) gene of 10-100 nucleotides comprising a fragment CC having a polymorphic site or a complement of the fragment. The TACR1 CC segment is useful for analysing a nucleic acid, by obtaining the nucleic acid from an individual, and determining the base occupying any one of the polymorphic sites in the segment. The nucleic acid is obtained from CC several individuals, and the base occupying one of the polymorphic sites is determined in each of the individuals, and further involves testing CC each of the individuals for the presence of a disease phenotype, and CC correlating the presence with the base. The TACR1 segment is useful for CC diagnosing and treating TACR1 ligand mediated diseases, such as asthma. CC The TACR1 segment is also useful in forensics, paternity testing, CC correlating polymorphisms with phenotypic traits, and genetic mapping of phenotypic traits. The TACR1 segment is useful in diagnosing and CC monitoring of diseases such as cancer, inflammation, heart disease, CC diseases of central nervous system, and susceptibility to infection to CC medicament for the treatment of the diseases. This polynucleotide correspond for the treatment of the diseases. This polynucleotide sequence represents a polymorphic region of the human neurokinin 1 CC receptor (TACR1) gene of the invention
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AGCCAGACTTCAGCAGC	48.0 76.5 ive	APPLICATION NUMBER: US/08/5 FILING DATE: 10/10/95 PRIOR APPLICATION DATA: APPLICATION NUMBER: 07/960, FILING DATE: 03/19/93 ANTORNEY/AGENT INFORMATION: NAME: Williams, ph.D., Kath REGISTRATION NUMBER: 34,380 REFERENCE/DOCKET NUMBER: 37 TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 345-9100 TELEFAX: (617) 345-9100 TELEFAX: (617) 345-9111 FORMATION FOR SEQ ID NO: 17: SEQUENCE CHARACTERISTICS: LENGTH: 17 bases TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: Synthetic RNA FEATURE: NAME/KEY: misc_feature LOCATION: 8 OTHER INFORMATION: N is 2'-541-950B-17	17 Application US/08541950B Application US/08541950B RMATION: Karn J, Gait MJ, Heaph NVENTION: VIRAL (HIV) G SEQUENCES: 26 ENCE ADDRESS: E: Banner & Witcoff, Lt One Financial Center, 4 One Financial Center, 4 Ill EADABLE FORM: EADABLE FORM: EADABLE FORM: CYPE: Diskette, 3.50 inc IBM PC compatible G SYSTEM: PC-DOS/MS-DOS		10 10 10 10 10 10 10 10 10 11
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TELEPAX: (617) 345-9111
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
STREET: One Financial Center, 45th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kd.
REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
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ADDRESSEE: Banner & Witcoff, Ltd.
                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                      APPLICANT: Karn J, OTITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Karn J, OT TITLE OF INVENTION: NUMBER OF SEQUENCES:
                               CURRENT APPLICATION DATA
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TELECOMMUNICATION INFORMATION:
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STATE: PL.
02111
TEAD/
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STATE: FL.
02111
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OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC
OPERATING SYSTEM:
                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WordPerfect 6.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc_feature LOCATION: '8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 0 FTT.ING DATE: 03/19/93
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
APPLICATION NUMBER: US/09/083,756A FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1490 AGCCAGACTTCAGCAGC 1506
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17 bases
                                                                                                                                                                                                                                                                   Karn J, Gait MJ, Heaphy S, Dingwall C
VENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
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                                                                                                        Diskette, 3.50 inch, 1.44 Mb storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       48.0%; Score 14.4; 76.5%; Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No. 6114109
GENERAL INFORMATION:
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Best Local Similarity
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TELEPHONE: (617) 345-9100
TELEFAX: (617) 345-9111
INFORMATION FOR SEQ ID NO: 17:
                                                                                                                                       TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTILLANT: KAYN J, Gait MJ, Heaphy S, Dingwall C
TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESS:
                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 3777/57347 (MRC-Ollax) TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathl
REGISTRATION NUMBER: 34,380
                                                                                                                                                                                                                                                                                                           FILING DATE: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: WordPerfect CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYFE:
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LENGTH: 17 bases
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                                 MOLECULE TYPE:
                                                                                                                       SEQUENCE CHARACTERISTICS:
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STATE: PL.
TD: 02111
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OPERATING SYSTEM: PC-DOS/MS-DOS
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LOCATION: 8
OTHER INFORMATION: N is 2'-deoxythymidine
                                                                TYPE: nucleic acid
STRANDEDNESS: singl
                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Williams, Ph.D., Kathleen
REGISTRATION NUMBER: 34,380
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   NAME/KEY:
                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Banner & Witcoff, Ltd.
STREET: One Financial Center, 45th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER:
                                                                                                     ENGTH:
                                                                                                                                                                       TELEPHONE:
                                                                                                                                                                                                                                                                           FILING DATE:
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                                                                                                     17 bases
misc_feature
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76.5%;
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Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                   1.44 Mb storage
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US-09-083-756A-23; Sequence 23; Application US/09083756A; Patent No. 6114109; GENERAL INFORMATION: APPLICANT: Karn J, Gait MJ, Heaph;
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US-08-541-950B-23
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                                                                                         RESULT 6
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                                                                                                                                                                                                                                             Query Match
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APPLICANT: Karn J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/
FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION UMBER: 07/960
APPLICATION UMBER: 07/960
FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 18 bases
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CURRENT APPLICATION DATA
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 345-9100
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                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: synthetic RNA
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STATE: PL.
To: 02111
                                                                                                                                                                                                            Local Similarity
nes 13; Conserv
                                                                                                                                                                                                                                                                                            OTHER INFORMATION: N is 4-thio-2'-deoxythymidine
                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature LOCATION: 8
                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Williams, Ph.D., REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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      Karn J, Gait MJ, Heaphy S, Dingwall C
                                                                                                                                                                                                            Conservative
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76.5%;
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                                                                                                                                                                                                        48.0%; Score 14.4; 176.5%; Pred. No. 3.1 cive 2; Mismatches
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Pred. No. 2
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                                           Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                 NUMBER OF SEQ ID NOS: 53
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 17
TYPE: RNA
ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/09325601 Patent No. 6573045
                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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                                                                                                                                                                                                                                     TITLE OF INVENTION: Methods and Kits for Discovery of RNA-Binding Compounds FILE REFERENCE: 3950/81235 CURRENT APPLICATION NUMBER: US/09/325,601 CURRENT FILING DATE: 1999-06-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (617) 345-91
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                       APPLICANT: Karn
APPLICANT: Prescott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/541,950
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen M.
REGISTRATION NUMBER: 34,380
REGISTRATION NUMBER: 34,380
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NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09,
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CIL.
STATE: Pa.
STATE: 02111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity nes 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: 8
OTHER INFORMATION: N is 4-thio-2'-deoxythymidine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE:
1490 AGCCAGACTTCAGCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1490 AGCCAGACTTCAGCAGC 1506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGCCAGANUUGAGCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           One Financial Center,
                                                    Conservative
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76.5%;
                                                                    46.0%;
76.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3777/57347 (MRC-011AX)
                                                                      Score 13.8;
Pred. No. 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.4; DB 1; Pred. No. 3.1;
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                                                    Mismatches
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                                                                                     DB 1;
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RESULT 9
US-09-083-756A-13
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                                                                                                                                                                                                                                                                                                Sequence 13, Application US/09083756A Patent No. 6114109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 5821046
GENERAL INFORMATION:
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FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/960,370
FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen
REGISTRATION NUMBER: 34,380
                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (617) 345-9111
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch,
COMPUTER: IRP Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
COMPUTER: IBM PC com
OPERATING SYSTEM: PC
                                                                                                                                                                    NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESSES: Banner & Witcoff, Ltd.
ADDRESSES: Banner & Witcoff, Ltd.
STREET: One Financial Center, 45th Floor
                                                                                                                                                                                                                                            APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: synthetic RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall CTITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                   C11.
STATE: Mr.
02111
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02111
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
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Similarity 76.5%;
                                                                                                                                                   Boston
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Pred. No. 3
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                                                                              1.44 Mb
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US-09-325-601-3
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                                                                                                                                                                                                                US-08-541-950B-18
                                                                                                                                                                                                                                 RESULT 11
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Best Local S
                                                                                                            Patent No. 5821046
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C
TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 53
SOPTWARE: PatentIn Ver. 2.1
SEQ ID NO 3
LENGTH: 18
TYPE: RNA
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                                                                                                                                                                                               Sequence 18, Application US/08541950B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Methods and Kits for Discovery of RNA-Binding Compounds FILE REFERENCE: 3950/81235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Karn
APPLICANT: Prescott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
ATTORNEY AGENT INFORMATION:
ANAME: Williams, Ph.D., Kathle.
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 3777
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/325,601
CURRENT FILING DATE: 1999-06-03
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LENGTH: 18 bases
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
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COMPUTER READABLE FORM:
   Cli.
STATE: MA
C2111
                             ADDAY: C. STREET: C. Boston
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                 Local
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                                                                                                                                                                                                                                                                                                                                                al Similarity
13; Conserv
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13; Conserv
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                                                               One Financial Center, 45th Floor
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                                                                                                                                                                                                                                                                                                                                                 Score 13.8; D
Pred. No. 3.9;
2; Mismatches
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Pred. No. 3
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                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 18;
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; LOCATION: 9
OTHER INFORMATION: N is 2'-deoxythymidine
US-08-541-950B-18
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                                APPLICATION NUMBER: US/08/541,95
ETILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/960,370
PILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
NAME: W1111ams, Ph.D., Kathleen
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 3777/57
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          atent No. 5821046
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                     STATE: MA
ZIP: 02111
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen M.
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 3777/57347 (MRC-O11AX)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07
FILING DATE: 03/19/93
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                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA
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                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WordPerfect 6.1
                                                                                                                                                                                                                                                                                                                                                                                                     STREET: One F
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE:
COMPUTER: II
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                                                                                                                                                                                                                                                                                                                                                                                                                         One Financial Center, 45th Floor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         42.7%; Score 12.8; Ilarity 76.5%; Pred. No. 5.6 Conservative 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Karn J, Gait MJ, Heaphy S, Dingwall C
ENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WordPerfect 6.1
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                                                                                                                                                                                                                                                                                                                           Diskette, 3.50 inch, 1.44 Mb storage
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                                                                                                                                                                                                                              US/08/541,950B
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                                                                                                                Kathleen M.
                                                                            3777/57347 (MRC-011AX)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.44 Mb storage
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                                   Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21, Application US/08541950B Patent No. 5821046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches 13;
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Best Local Similarity
                                                                                                                                                                                                                                                            TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08,
FILLING DATE: 10/10/95
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 07/9
FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Banner & Witcoff, Ltd.
                                                                                                                                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall CTITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: 10 OTHER INFORMATION: N is 2'-deoxythymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
                                                  Local
                                                                                                               LOCATION: 9 -
OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
                                                                                                                                           NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                            NAME: Williams, Ph.D., REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                                                                                                                                                                                                           ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1490 AGCCAGACTTCAGCAGC 1506
   1490 AGCCAGACTTCAGCAGC 1506
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                                                    Similarity
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                                                                                                                                                                                                                                                17 bases
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                                                                                                                                                                                synthetic RNA
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                                                  42.7%;
76.5%;
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                                                                                                                                                                                                                                                                                                                                                                           Kathleen M.
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                                   Score 12.8; D
Pred. No. 5.6;
1; Mismatches
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Pred. No. 5.
                                                                 DB 1; Length 17;
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Sequence 22, Appr-
No. 5821046
                                                                                                                                                                                                                                                Sequence 18, Application US/09083756A Patent No. 6114109
                                                                                                                                                                                                                               Patent No. 6114109
GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO: 22:
                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/541,9
FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/960,370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: synthetic RNA FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                     ADDATE CALL STREET: CALL BOSTON
                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen M.
REGISTRATION NUMBER: 34,380
REGISTRATION NUMBER: 34,380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA
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STATE: PL.
02111
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STATE: Pr.
To: 02111
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STRANDEDNESS: singl
TOPOLOGY: linear
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION: 10 OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM:
SOFTWARE: WordPer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER:
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                                                                                                                                                                                                                                                                                                                                                                                                1490 AGCCAGACTTCAGCAGC 1506
                                                                                                                                                                                                                                                                                                                                                                                                                                     13;
                                                                                                                                                                                                                                                                                                                                                              1 AGCCAGAUUNGAGCAGC 17
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                                                                                                                     Banner & Witcoff, Ltd.
Pe Financial Center, 45th Floor
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IBM PC compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  42.7%;
76.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 12.8;
Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No. 6114109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19, Application US/09083756A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                       TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Karn J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO:
                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                          FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen M.
REGISTRATION NUMBER: 34,380
                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WORDERFECT 6.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                      STATE: MA
ZIP: 02111
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 17 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 345-9100
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APPLICATION NUMBER:
FILING DATE:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09,
                                                                                                                         REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: synthetic RNA
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STREET: MA
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LOCATION: 9
OTHER INFORMATION: N is 2'-deoxythymidine
                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Williams, Ph.D., Kathi
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE
TYPE: nucleic acid STRANDEDNESS: singl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                       TELEPHONE:
                                                                                                                                                                                                                                    APPLICATION NUMBER:
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                                      ENGTH:
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                                      17 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Karn J, Gait MJ, Heaphy S, Dingwall C
/ENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                          (617)
                                                                                                                                                                                                                                                                                                                                                                            Diskette, 3.50 inch, 1.44 Mb
                                                                                                          345-9100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Score 12.8; D; Pred. No. 5.6; 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3777/57347 (MRC-011AX)
                                                                                                                                             3777/57347
                                                                                                                                               (MRC-011AX)
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                                                                                                                                                                                                                                                                                                                                                                              storage
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TOPOLOGY: linear MOLECULE TYPE: synthetic RNA FEATURE:

NAME/KEY: misc_feature

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RESULT 18
US-09-083-756A-22
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                                                                                                                                                                                                                ; NAME/KEY: misc_feature
LOCATION: 9
; OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
US-09-083-756A-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-083-756A-19
                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATE:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleer
NAME: Williams, Ph.D., Kathleer
NAME: Williams, Ph.D., Kathleer
NAME: Williams, Ph.D., Kathleer
NAME: 34,380
REGISTRATION UMBER: 3777/
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
(617) 345-9110
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Matches
                                                                                                                                         Query Match 42. 
Best Local Similarity 76. 
Matches 13; Conservative
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/083,756A FILING DATE: PRIOR APPLICATION DATA: APPLICATION NUMBER: 08/541,950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
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TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb
                                                                                                                                                                                                                                                                                                          NOLECULE TYPE:
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STATE: PL.
STATE: 02111
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SOFTWARE: WordPer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: 10
OTHER INFORMATION: N is 2'-deoxythymidine
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les 13; Conserv
                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                               ENGTH:
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                                                                       1 AGCCAGAUNUGAGCAGC 17
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                                                                                                                                                                                                                                                                                                                                                                              17 bases
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SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                        synthetic RNA
                                                                                                                                                           42.7%;
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76.5%; Pred. No. 5.6;
tive 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kathleen M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3777/57347 (MRC-Ollax)
                                                                                                                                                             Score 12.8;
Pred. No. 5.
                                                                                                                                          Mismatches
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                                                                                                                                                                            DB 1; Length 17;
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RESULT 19
US-08-363-240A-47/c
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                                                                                                                                                                                                                                                                           Patent No.
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (617) 345-9100
TELEFAX: (617) 345-9111
INFORMATION FOR SEQ ID NO: 22
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09,
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MEDIUM TYPE: Diskette, 3.50 inch,

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                         APPLICANT: Pape, Michael TITLE OF INVENTION: METH
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                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                          TITLE OF INVENTION:
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TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
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STATE: "".
Co. 02111
                                        ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
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LOCATION: 10
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STRANDEDNESS: single
TOPOLOGY: linear
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COUNTRY:
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                                                                                                            OF SEQUENCES:
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           Los Angeles
California
                                                                                                                                                                                                                                                                                        Application US/08363240A
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Bisgaier, Charles
                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        synthetic RNA
                                                                                                                          METHOD AND REAGENT FOR PREVENTION, INHIBITION OF PROGRESSION AND REGRESSION OF VASCULAR DISEASES
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76.5%;
                                                                                                            1243
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                                                             Street
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Pred. No. 5.
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Patent No. 5
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Best Local Similarity 86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Petry, Douglas A.
REGISTRATION NUMBER: 35,321
REFERENCE/DOCKET NUMBER: 8765
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US
FILING DATE: December
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIE: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                    STREET: 340 K
CITY: Nutley
STATE: New Je:
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                                                                                      APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                        COUNTRY: U
ZIP: 07110
                                                                                                                                                                                                                                                                              ADDRESSEE:
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5567809
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                                                                                                                                                                                                                                                                                 340 Kingsland
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Begovich, Aum -.
Teodorica L.
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                                                                                                                                                                                                                                                                             Hoffmann-La Roche Inc
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                                                                                                                                                                                                                                                                                                                                                                                                                            Raymond
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Pred. No. 7
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Query Match
Best Local Similarity
Matches 12; Conserve
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                                                                                                                                                                                                                                                                                                                                 US-09-508-753B-25/c
                                                                                                                                                                                                                                                                                                                                                   RESULT 22
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APPLICANT: Indue, Takakazu
APPLICANT: Indue, Takakazu
TITLE OF INVENTION: Method of Amplifying DNA Fragment, Apparatus for Amplifying DNA Fr
TITLE OF INVENTION: agment, Method of Assaying Microorganisms, Method of Analyzing Mic
TITLE OF INVENTION: nisms and Method of Assaying Contaminant
FILE REPERENCE: 9982-7
CURRENT APPLICATION NUMBER: US/09/281,418
CURRENT FILING DATE: 1999-03-31
EARLIER APPLICATION NUMBER: JP/1998/87651
EARLIER FILING DATE: 1998-03-31
                                                                                                                                                                   APPLICANT: AKIRA SHIMAMOTO
APPLICANT: Yasuhiro FURUICI
APPLICANT: Yuko SHIBATA
APPLICANT: Hiroko FUNAKI
APPLICANT: Eiji OHARA
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                                                                                                                                                                                                                                                                      Patent No. 6544736 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                        Sequence 25, Application US/09508753B Patent No. 6544736
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 65
NUMBER OF SEQ ID NOS:
SEQ ID NO 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                  CURRENT APPLICATION NUMBER: US/09/508,753B
CURRENT FILING DATE: 2000-06-16
PRIOR APPLICATION NUMBER: JP 9/270324
PRIOR FILING DATE: 1997-09-18
                                                                                                                 APPLICANT: Masanori WATAHIKI
TITLE OF INVENTION: Method for Synthesizing
FILE REFERENCE: 00162/HG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EARLIER APPLICATION NUMBER: JP/1999/69694
EARLIER FILING DATE: 1999-03-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Primer
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Yuko SHIBATA
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Pred. No. 16;
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Pred. No. 8.7;
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                                                                                                                                     cDNA from mRNA sample
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RESULT 24
US-09-721-777-18
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Best Local Similarity
'---hes 9; Conserv:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-263-790-36
CURRENT APPLICATION NUMBER: US/09/721,777
CURRENT FILING DATE: 2000-11-27
NUMBER OF SEQ ID NOS: 20
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 18
EQ ID NO 18
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Rest Local Similarity 100.
9; Conservative
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-09-263-790-36
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SEQ ID NO 36
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APPLICANT: Nirmal Kumar PATRA et al.

APPLICANT: OF INVENTION: JAL PALLAVI, WATER LOGGING TOLERANT CYMBOPOGON WINTERIANUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 36, Application US/09263790 Patent No. PP12997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/263,790
CURRENT FILING DATE: 1999-03-05
NUMBER OF SEQ ID NOS: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 10
TYPE: DNA
ORGANISM: Artificial
FEATURE:
                                                                                                                                                                                                                             APPLICANT:
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INVENTION: Mint Plant Named Saksham
                                                                                                                                                                                                                                                                                                                                                                                  PP13279
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                                                                                                                                                                                    Singh, Anil Kumar
Patra, Nirmal Kumar
Bahl, Janak Raj
                                                                                                                                                                                                                                                                                          Shasany, Ajic ,
Dhawan, Sunita
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                                                                                                                                                                                                                                          Naqvi, Ali Arif
Dhawan, Om Parkash
                                                                                                                                                                                                                                                                             Darokar, Mahendra Pandurang
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100.0%; Pred. No. 16;
cive 0; Mismatches
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Pred. No. 2
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Query Match
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                                                                                                          US-08-545-253A-20
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Best Local Similarity 90.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                    TELEX: 575102
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                           REGISTRATION NUMBER: 31,665
REFERENCE/DOCKET NUMBER: 50
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 881-3140
                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenneth D. Sibley
STREET: Post Office Drawer 34009
                                                                                                                    TOPOLOGY: 15 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Sibley, Kenneth D.
                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                  STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                 TELEPHONE: (915) 881-3175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Charlotte
                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
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                        1488 GAAGCCAGAC 1497
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IVENTION: METHODS FOR WITHIN FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                U.S.A.
                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Henry V. Amers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Grattapaglia, Dario
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90.0%;
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                                                               Score 8.4;
Pred. No.
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Pred. No.
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RESULT 26 US-08-719-337-20

Sequence 20, Applion Patent No. 6054634

Application US/08719337

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RESULT 27
US-09-255-432-6
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CLASSIFICATION: 047

CLASSIFICATION DATA:
APPLICATION NUMBER: 08/184,567

FILING DATE: 21-JAN-1994

ATTORNSY/AGENT INFORMATION:
NAME: Sibley, Kenneth D.
REGISTRATION NUMBER: 31,665

REGISTRATION NUMBER: 5051-2

TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
TOTREPHONE: (919) 881-3140
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; MOLECULE TYPE:
US-08-719-337-20
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Best Local Similarity
Matches 9; Conserva
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                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: POST CITY: Charlotte
STATE: No. 6054634th Carolina
TOWNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: METHODS FOR WITHIN FAMILY SELECTION IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 575102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (919) 881-3175
                                                                                                                           ADDRESSEE: Judy C. Jarecki-Black, Ph.D.
ADDRESSEE: Dority & Manning, P.A.
STREET: 700 E. No. 6258537th Street, Suite 15
CITY: Greenville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: single
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             COMPUTER: IBM con
OPERATING SYSTEM:
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                                                                                            COUNTRY:
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                                                                                                             South Carolina
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Post Office Drawer 34009
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Sederoff,
                                                                                              USA
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IBM compatible
SYSTEM: MS Dos; Windows
WordPerfect 6.1
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                                             Diskette,
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                                             3.50 inch, 1.44 Mb storage
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Pred. No. :
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                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: AFFILING DATE: 21 June:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (864) 233-7342 INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ANTI-SENSE: NO
ANTI-SENSE: NO
ORIGINAL SOURCE: Operon Technologies (Alemeda,
IMMEDIATE SOURCE: Operon Technologies
posttion in Genome: No. 6258537 Applicable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: Filed Her
                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                         COMPUTER: IBM Compatible w/
OPERATING SYSTEM: Microsoft
SOFTWARE: Microsoft word 97
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:

ADDRESSEE: D. Peter Hochberg Co., L.P.A.
STREET: The Baker Building - Sixth Floor 1940 East 6th Street
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: William Mitchell Molyneux TITLE OF INVENTION: Mosquito and/or Flea Control NUMBER OF SEQUENCES: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
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DESCRIPTION:
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REFERENCE/DOCKET NUMBER: CX
TELECOMMUNICATION INFORMATION:
TELEPHONE: (864) 271-1592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA: C1
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                     STREET: The STATE: Ohio
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                STRANDEDNESS:
                                                                                                                                                                        APPLICATION NUMBER: US
FILING DATE: June 19,
                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: Linear
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                                                   ENGTH:
                                                                                                                                                           CLASSIFICATION:
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                               Nucleic Acid
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Linear
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                                                                                                                                                                                                                                                                                  Diskette,
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No. 6258537 Applicable
                                                                                                                        AU PO 0605
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                                                                                                      1996
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Query Match
Best Local Similarity
 28.0%;
 Score 8.4; D
Pred. No. 20;
           DB 1;
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FILE REFERENCE: 00162/HG
CURRENT APPLICATION NUMBER: US/09/508,753B
CURRENT FILING DATE: 2000-06-16
PRIOR APPLICATION NUMBER: JP 9/270324
PRIOR FILING DATE: 1997-09-18
NUMBER OF SEQ ID NOS: 472
SEQ ID NO 63
LENGTH: 10
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 30
US-09-508-753B-63/c
                                     ; OTHER INFORMATION: Description of Artificial Sequence: Primer US-09-508-753B-63
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CURRENT APPLICATION NUMBER: US/09/508,753B
CURRENT FILING DATE: 2000-06-16
PRIOR APPLICATION NUMBER: JP 9/270324
PRIOR FILING DATE: 1997-09-18
NUMBER OF SEQ ID NOS: 472
SEQ ID NO 28
LENGTH: 10
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Best Local Similarity
Matches 9; Conserv
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Best Local Similarity
Matches 9; Conserv
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                APPLICANT: Eiji OHARA
APPLICANT: MaBanori WATAHIKI
TITLE OF INVENTION: Method for Synthesizing cDNA from mRNA sample
                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Akira SHIMAMOTO
APPLICANT: Yasuhiro FURUICHI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Akira SHIMAMOTO APPLICANT: Yasuhiro FURUI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Masanori WATAHIKI
FITLE OF INVENTION: Method for Synthesizing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: Primer
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                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
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Yuko SHIBATA
                                                                                                                                                                                                                                                                                                                            Hiroko FUNAKI
                                                                                                                                                                                                                                                                                                                                           Yuko SHIBATA
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Pred. No. 20;
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Pred. No. 20;
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Length 10;
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                                                      Sequence 6, Application US/09758073 Patent No. 6610487 GENERAL INFORMATION:
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Best Local Similarity
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ADDRESSEE: The Webb Law Firm
STREET: 700 Koppers Building, 436 Seventh Avenue
           APPLICANT: Keinath, TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 412-471-8815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/EP
FILING DATE: 19-FEB-1996
APPLICATION NUMBER: 952003
FILING DATE: 17-FEB-1995
APPLICATION NUMBER: 952019
FILING DATE: 14-UUL-1995
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: SCHOENMAKERS, H.F.D.M.
TITLE OF INVENTION: MULTIPLE-TUMOR ABERRENT GROWTH
TITLE OF INVENTION: GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 70:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Johnson, Barbara E REGISTRATION NUMBER: 31,
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et al.
Method of Diagnosing Gummy Stem Blight in
Plants Using a Polymerase Chain Reaction Assay
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TUMBER OF SEQUENCES:

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US-08-173-489C-342/c
; Sequence 342, Application US/08173489C
; Patent No. 5861244
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Best Local Similarity
"hes 9; Conserv
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                                                                                                                                                                                    Patent No. 5861244
GENERAL INFORMATION:
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MOLECULE TYPE:
DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 16-MAR-1998
ATTORNEY/AGENT INFORMATION:
NAME: Judy C. Jarecki-Black, Ph.D.
REGISTRATION NUMBER: P44,170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (864) 271-1592
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                                                                                        NUMBER OF SEQUENCES: 3
                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORIGINAL SOURCE: Operon Technologies (Alemeda, IMMEDIATE SOURCE: Operon Technologies POSITION IN GENOME: No. 6610487 Applicable
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Dority & Manning, P.A. STREET: 700 E. No. 6610487th Street, Suite CITY: Greenville
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OPERATING SYSTEM:
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                           CITY: NEW YORK
STATE: NEW YORK
                                                           STREET:
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                                                                          ADDRESSEE:
                COUNTRY:
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HEPBURN,
                USA
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Oligonucleotide Primer
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GENETIC SEQUENCE ASSAY USING DNA
TRIPLE-STRAND FORMATION.
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Pred. No. 20;
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                                                                                                              ; OTHER INFORMATION: Symthetic oligonucleotide primer US-09-862-847-15
                                                                                                                                                                                                                                  SOFTWARE: PatentIn version 3.1 SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15, Applica Patent No. 6593111 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                  TITLE OF INVENTION: DIRECTION ASSEMBLY OF LARGE VIRAL GENOMES AND CHROMOSOMES FILE REFERENCE: 5470.270
CURRENT APPLICATION NUMBER: US/09/862,847
CURRENT FILING DATE: 2001-05-21
PRIOR APPLICATION NUMBER: US 60/206,537
PRIOR FILING DATE: 2000-05-21
PRIOR PRIOR FILING DATE: 2000-05-21
PRIOR APPLICATION NUMBER: US 60/285,320
PRIOR FILING DATE: 2001-04-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Baric, Ralph S. APPLICANT: Boyd, Youn:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (attorney) (:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                ENGTH: 11
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MOLECULE TYPE: other nucleic acid

DESCRIPTION: third strand derived from P. cepacea

DESCRIPTION: 16s region in Seq ID No. 5861244341
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LENGTH: 11 bases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inch, 1.44Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Handelman, Joseph H.
REGISTRATION NUMBER: 26,179
REFERENCE/DOCKET NUMBER: U9518-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
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1484 CCAAGAAGCC 1493
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                                        Conservative
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Pred. No. 21;
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RESULT 36
US-09-041-675-19/c
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                                                                                                              GENERAL INFORMATION:
                                                                                                                                Sequence 19, Application US/09041675A Patent No. 6100032
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APPLICANT: Kern, Scott
APPLICANT: Zawel, Leigh
APPLICANT: Dai, Jia Le
APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Human SMAD3 and SMAD4 are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,714
REFERRNCE/DOCKET NUMBER: D-59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/651-5325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: 713/651-532:
TELEPA: 73/651-5246
TIMPORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
NOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "oligonucleotide"
"VPOTHETICAL: YES
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APPLICANT:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPANIAN
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/859,954
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PRIOR APPLICATION DATA:
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APPLICANT: Hardin, Paul E.
TITLE OF INVENTION: Design and Optimized Primer Librar
TITLE OF INVENTION: Gene Sequencing and Method Thereof
NUMBER OF SEQUENCES: 566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
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1301 McKinney, Suite 5100
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GENERAL INFORMATION:
APPLICANT: Kern, Scott
APPLICANT: Dai, Jia Le
APPLICANT: Dai, Jia Le
APPLICANT: Dai, Jia Le
APPLICANT: Vogelatein, Bert
APPLICANT: Vogelatein, Bert
APPLICANT: Vogelatein, Bert
APPLICANT: Winzler, Kenneth
TITLE OF INVENTION: Human SMAD3 and SMAD4 are
TITLE OF INVENTION: sequence-specific transcription activators
FILE REFERENCE: 01107.74098
CURRENT APPLICATION NUMBER: US/09/041,675A
CURRENT APPLICATION SUMMER: US/09/041,675A
CURRENT FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 24
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US-09-989-789-455/c
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GENERAL INFORMATION:
APPLICANT: LIU, QiANG
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/09/989,789
CURRENT FILING DATE: 2002-03-25
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
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CURRENT FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 19
                                                                                                                                                                                Sequence 455, Application US/09989789
Patent No. 6588746
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 8
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
OTHER INFORMATION: synthetic random oligonucleotides selected
OTHER INFORMATION: binding to human SMAD3 or human SMAD4
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TYPE: DNA
ORGANISM: Artificial Sequence
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100.0%; Pr
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; OTHER INFORMATION: Description of Artificial Sequence: example target ; OTHER INFORMATION: DNA US-09-989-789-455
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APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
TILLE REFERENCE: 835-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/09/989,789
CURRENT FILING DATE: 2002-03-25
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 456
LENGTH: 9
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: example target OTHER INFORMATION: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
               ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                     APPLICANT: Jerry W. Shay
APPLICANT: Woodring E. Wright
APPLICANT: Elizabeth Blackburn
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF CONDITIONS
TITLE OF INVENTION: RELATED TO TELOMERE LENGTH AND/OR
                                                                                                                                                                                                                       TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                             STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                    STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 100
les 8; Conservative
                                                                                      COUNTRY: U.S.A.
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Compatible
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100.0%; Pr
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0; Mismatches
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; Pred. No. 1.2e+02;
0; Mismatches 0;
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Query Match
Best Local Similarity
Matches 8; Conserv:
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; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 4
; LENGTH: 10
; TYPE: DNA
; ORGANISM: Mycobacterium tuberculosis
US-08-997-897-4
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Best Local Similarity
Watches 8; Conservi
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                                                                                          RESULT 42
US-09-156-836B-4
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                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/08997897C
Patent No. 6114514
GENERAL INFORMATION:
APPLICANT: SRIVASTAVA, RANJANA
APPLICANT: KUMAR, DEEPAK
APPLICANT: KUMAR, DEEPAK
APPLICANT: SRIVASTAVA, BRAHM SHANKER
TITLE OF INVENTION: MYCOBACTERIUM TUBERCULOSIS SPECIFIC DNA FRAGMENT
                                            Sequence 4, Application US/09156836B Patent No. 6242585 GENERAL INFORMATION:
 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: u011469-7
CURRENT APPLICATION NUMBER: US/08/997,897C
CURRENT FILING DATE: 1997-12-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 07/882,431
FILING DATE: May 13, 1992
APPLICATION NUMBER: 08/018,766
FILING DATE: March 24, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 202//
TELECHMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 489-1600
                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: May 13,
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
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TOPOLOGY: 1i
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM:
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                                                                                                                                                       AAGAAGCC 10
Srivastava, Ranjana
Kumar, Deepak
Srivastava, Brahm Sh
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                                                                                                                                                                                                                  Conservative
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May 13, 1992
UMBER: 08/038,766
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5. 23;
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RESULT 43
US-08-464-011B-9/c
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9, Application US/08464011B Patent No. 6368789 GENERAL INFORMATION:
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SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/156,836B
CURRENT FILING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 08/997,897
PRIOR FILING DATE: 1997-12-24
NUMBER OF SEQ ID NOS: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: MYCUBAL 011876 4
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TYPE: DNA
ORGANISM: Mycobacterium tuberculosis
                                                     TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 9:
                                                                                                                                           APPLICATION NUMBER: 07/882,438
FILING DATE: Way 13, 1992
APPLICATION NUMBER: 08/038,766
FILING DATE: March 24, 1993
APPLICATION NUMBER: 08/66,952
FILING DATE: May 13, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Warthurg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 202/045
                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/464,011B
FILING DATE: 05-Jun-1995
CLASSIFICATION: -Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOPTMARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 61
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Woodring E. Wright
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF CONDITIONS
RELATED TO TELOMERE LENGTH AND/OR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1486 AAGAAGCC 1493
                               SEQUENCE CHARACTERISTICS:
                                                                                                        TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Michael D. West
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LENGTH: 10
TYPE: nucle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                      TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MYCOBACTERIUM TUBERCULOSIS SPECIFIC DNA FRAGMENT
                                                                                                          (213) 489-1600
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Pred. No. 23;
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Search completed: April 15, 2004, 16:36:47 Job time : 0.001 secs

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Query Match
Best Local Similarity
""" hes 8; Conserv:
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                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence:Synthetic oligonucleotide US-09-336-946B-15
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US-09-336-946B-15
                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/336,946B; CURRENT FILING DATE: 1999-06-21
PRIOR APPLICATION NUMBER: 60/095229
PRIOR FILING DATE: 1998-08-04
NUMBER OF SEQ ID NOS: 74
SOFTWARE: Microsoft Office 97
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15, Application US/09336946B Patent No. 6479731
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Best Local S
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                                                                                                                                                                                                                                                                                                                                     APPLICANT: Bryan, Gregory
APPLICANT: E. I. du Pont de Nemours and Company
TITLE OF INVENTION: A Pi-ta GENE CONFERRING DISEASE RESISTANCE TO PLANTS
FILE REFERENCE: BB-1136
                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Valent, Barbara S.
                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                               FEATURE:
                                                                                                                                                                                                             LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 100
nes 8; Conservative
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                            1490 AGCCAGAC 1497
2 AGCCAGAC 9
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                                                             Conservative
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100.0%;
                                                         26.7%; Score 8; DB 1
100.0%; Pred. No. 23;
tive 0; Mismatches
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o. 23;
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9, Appli 185, App 1011, Ap

344, 3, Appli 3, Appli 5, Appli 6, Appli 147, App

455,

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 44 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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30.420 Million cell updates/sec
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   US-09-954-556-65
US-09-954-556-67
US-09-954-556-67
US-09-954-556-67
US-10-307-005-1800
US-10-004-551-43
US-10-004-551-43
US-09-504-231A-1451
US-09-504-231A-1451
US-09-274-553D-1435
US-09-274-553D-1435
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US-09-274-553D-1435
US-09-274-553D-1435
US-09-274-553D-144
US-09-353-145-1647
US-10-106-799-13
US-10-106-799-14
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US-09-982-250-1651
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US-10-330-627-110
US-10-330-627-110
US-09-9862-847-15
US-09-9862-847-15
US-09-9862-847-15
US-09-986-2848-1661
US-09-989-789-456
US-09-989-789-456
US-09-990-186-456
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Sequence 65, Appl
Sequence 67, Appl
Sequence 1799, Ap
Sequence 1800, Ap
Sequence 40, Appl
Sequence 41, Appl
Sequence 1451, Ap
Sequence 1451, Ap
Sequence 1135, Ap
Sequence 1137, Ap
Sequence 1137, Ap
Sequence 11, Appl
Sequence 11, Appl
Sequence 17, Appl
Sequence 14, Appl
Sequence 14, Appl
Sequence 1647, Appl
Sequence 1647, Appl
Sequence 1671, Appl
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Sequence 1107, Appl
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Sequence 455, Appl
Sequence 16, Appl
Sequence 455, Appl
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APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper

ITITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPRES
FILE REFERENCE: RTS-0250
CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NO.66
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Antisense Oligonucleotide
US-09-954-556-66
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US-09-954-556-65/c
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SEQ ID NO 65
LENGTH: 20
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APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
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US-10-277-494-147
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; Sequence 67, Application US/09954556
; Publication No. US20030078219A1
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CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 67
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                         Query Match
Best Local Similarity
                                                                               Matches
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Publication No. US20030236208A1
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APPLICANT: Howard B. Gamper
APPLICANT: Michael C. Rice
APPLICANT: Michael C. Rice
APPLICANT: Jungsup Kim
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations in Plants
TITLE OF INVENTION: Using Modified Single Stranded Oligonucleotides
FILE REFERENCE: Napro/009 PCT
CURRENT APPLICATION NUMBER: US/10/307,005
CURRENT APPLICATION NUMBER: PC7/US01/17672
                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS:
SOFTWARE: Friedman mac
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PRIOR APPLICATION NUMBER: US 60/208,538

PRIOR FILING DATE: 2000-06-01

PRIOR APPLICATION NUMBER: US 60/244,989

PRIOR FILING DATE: 2000-10-30

PRIOR APPLICATION NUMBER: US 09/818,875

PRIOR FILING DATE: 2001-03-27

PRIOR FILING DATE: 2001-03-27
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APPLICANT: SCOTT COOPER
TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2
FILE REFERENCE: RTS-0250
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APPLICANT: Susan M. Frei
APPLICANT: Scott Cooper
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                                    1485 CAAGAAGCCAGACTTCA 1501
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100.0%; Pred. No. 0.
Live 0; Mismatches
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Pred. No. 4
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Publication No. US20030004310A1
GENERAL INFORMATION:
APPLICANT: SHIMKETS, RICHARD A
APPLICANT: FERNANDES, ELMA
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SEQ ID NO 1800
LENGTH: 17
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CURRENT FILING DATE: 2001-12-05
PRIOR APPLICATION NUMBER: 09/635,949
PRIOR FILING DATE: 2000-08-10
NUMBER OF SEQ ID NOS: 110
SOFTWARE: Patentin Ver. 2.1
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PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 2717
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TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES ENCODED THEREBY
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APPLICANT: Howard B. Gamper
APPLICANT: Michael C. Rice
APPLICANT: Jungs:p Kim
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Pred. No. 4.8;
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Pred. No. 5
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US-09-504-231A-1451/c
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 43
LENGTH: 18
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SEQ ID NO 1451
LENGTH: 14
                                                                                      Query Match
Best Local Similarity
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Matches 15; Conserv
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                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 1999-02-24
PRIOR APPLICATION NUMBER: 60/100,842
PRIOR FILING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 60/083,217
PRIOR FILING DATE: 1998-04-27
NUMBER OF SEQ ID NOS: 3242
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CURRENT APPLICATION NUMBER: US/10/004,551
CURRENT FILING DATE: 2001-12-05
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PRIOR FILING DATE: 2000-08-10
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PRIOR FILING DATE: 1999-03-23
PRIOR APPLICATION NUMBER: 09/257,608
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0. US20020013458A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Roberts, be
Roberts, be
Pamela
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                                   CTTCAGCAGCCA 1508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         McSwiggen, Jan
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                                                                       Conservative
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88.2%;
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                                                                                       Score 12; DB 1; ; Pred. No. 7.3;
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Pred. No. 5
                                                                       Mismatches
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US-09-504-231A-1435/c
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Best Local Similarity
Matches 12; Conservi
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; OTHER INFORMATION: Description of Artificial Sequence: US-09-504-231A-1435
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 1451
                                                                                                                             SOFTWARE: PatentIn
SEQ ID NO 1435
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APPLICANT: Macejak, Dennis
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES OR CONDITIONS RELATEL
TITLE OF INVENTION: HEPATITIS C VIRUS INFECTION
FILE REFERENCE: rpi 247/282
FILE REFERENCE: rpi 247/282
                                                                                                                                                                                                                                     PRIOR FILING DATE: 1999-03-23
PRIOR APPLICATION NUMBER: 09/257,608
PRIOR FILING DATE: 1999-02-24
PRIOR APPLICATION NUMBER: 60/100,842
PRIOR FILING DATE: 1998-09-18
                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: xpi 247/282
CURRENT APPLICATION NUMBER: US/09/504,231A
CURRENT FILING DATE: 2000-02-15
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PRIOR APPLICATION NUMBER: 60/100,842
PRIOR FILING DATE: 1998-09-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: McSwiggen, Jan
APPLICANT: Roberts, Beth
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                                                                                                                                                                                               PRIOR FILING DATE: 1998-04-27
                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/083,217
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Blatt,
APPLICANT: McSwi
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                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                         PPLICANT: Macejak, Dennis
ITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES OR CONDITIONS RELATED
ITLE OF INVENTION: HEPATITIS C VIRUS INFECTION
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Pred. No.
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7.3;
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Query Match
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                                                              ; ORGANISM: Homo sapiens US-10-033-145-1327
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PRIOR FILING DATE: 1999-02-24
PRIOR APPLICATION NUMBER: 60/100,842
PRIOR FILING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 60/083,217
PRIOR FILING DATE: 1998-04-27
NUMBER OF SEQ ID NOS: 3148
                                                                                                                           NUMBER OF SEQ ID NOS: 2137
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1327
LENGTH: 10
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Publication No. US20020151515A1
GENERAL INFORMATION:
Query Match
Best Local Similarity
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APPLICANT: GENZYME CORPORATION
APPLICANT: SHANKARA, SRINIVAS
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
FILE REFERENCE: GA0201C
CURRENT APPLICATION NUMBER: US/10/033,145
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
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APPLICANT: McSwiggen, James
APPLICANT: Roberts, Beth
APPLICANT: Roberts, Beth
APPLICANT: Pavco, Pamela
APPLICANT: Macejak, Dennis
APPLICANT: Macejak, Dennis
TITLE OF INVENTION: ENZWATIC NUCLEIC ACID TREATMENT OF DISEASES OR CONDITIONS RELATE
TITLE OF INVENTION: ENZWATIC VIRUS INFECTION
FILE REFERENCE: Tpi 247/282
CURRENT APPLICATION NUMBER: US/09/274,553D
CURRENT APPLICATION NUMBER: US/09/274,553D
CURRENT FILING DATE: 1999-03-23
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TYPE: RNA
                                                                                                          TYPE: DNA
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33.3%;
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85.7%; Pred. No. 1:
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Pred. No. 11;
  Score 10;
Pred. No.
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  DB 1;
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Query Match
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RESULT 15
US-10-240-580-10/c
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US-10-106-799-13
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OTHER INFORMATION: primer_bind
US-09-823-887C-17
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                                                                                                                                                                                                                                                                                                  Sequence 13, Application US/10106799
Publication No. US20030140379A1
GENERAL INFORMATION:
APPLICANT: Council of Scientific and Industrial Research
TITLE OF INVENTION: No. US20030140379A1e1 DNA sequence in plants Caragana jubata with
TITLE OF INVENTION: method thereof
FILE REFERENCE: US 673
CURRENT APPLICATION NUMBER: US/10/106,799
CURRENT FILING DATE: 2002-10-31
NUMBER OF SEQ ID NOS: 32
SOFTWARE: Patentin version 3.1
150 ID NO 13
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 17
LENGTH: 13
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APPLICANT: Lal, Lakhvir
APPLICANT: Lal, Lakhvir
TITLE OF INVENTION: Cloning of No. US20030180723A
TITLE OF INVENTION: Dormancy in the Apical Buds
FILE REFERENCE: H053916.0001US0
CURRENT APPLICATION NUMBER: US/09/823,887C
CURRENT FILING DATE: 2002-04-23
                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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Sequence 10, Application US/10240580 Publication No. US20030180716A1

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RESULT 16
US-08-935-377-14
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Publication No. US20030133917A1
GENERAL INFORMATION:
APPLICANT: Zauderer, Maurice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 31.3
Best Local Similarity 90.9
Matches 10; Conservative
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SEQ ID NO 10
LENGTH: 12
TYPE: DNA
Query Match
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CURRENT FILING DATE: 2002-09-30
PRIOR APPLICATION NUMBER: PCT/JP01/02516
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: JP 2000-99482
PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: INOUE, Takakazu

TITLE OF INVENTION: METHOD AND APPARATUS FOR MICROORGANISM DISCRIMINATION, METHOD OF

TITLE OF INVENTION: DATABASE FOR MICROORGANISM DISCRIMINATION, AND MICROORGANISM DI

TITLE OF INVENTION: PROGRAM AND RECORDING MEDIUM ON WHICH SUCH PROGRAM IS RECORDED

FILE REFERENCE: 9982-24
                                                                                                                                                                                                                                 CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K
REGISTRATION NUMBER: 36,688
REGERENCE/DOCKET NUMBER: 1821
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                  TELEFAX: (202) 371-2540
NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
OTHER INFORMATION: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: T Cells Specific for Target Antigens and TITLE OF INVENTION: Vaccines Based Thereon
                                                                 MOLECULE TYPE:
                                                                                                   STRANDEDNESS:
                                                                                                                           TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/0
FILING DATE: 22-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C STREET: 1100 New York Avenue, N.W., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Washington
                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1484 CCAAGAAGCCA 1494
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Pred. No. 15;
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30.0%;

Score 9;

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1;

Length 10;

GENERAL INFORMATION:

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APPLICANT: Zauderer, Maurice
APPLICANT: Zauderer, Maurice
TITLE OF INVENTION: Methods for Producing Recombinant Libraries in Vaccinia Virus
FILE REFERENCE: 1821.0010001
CURRENT APPLICATION UNMBER: US/09/822,250
CURRENT FILING DATE: 2001-04-02
PRIOR APPLICATION NUMBER: US 08/935,377
PRIOR FILING DATE: 1997-09-22
NUMBER OF SEQ ID NOS: 37
SOFTWARE: PatentIn version 3.0
SEQ ID NO 14
LENGTH: 10
TYPE: DNA
ORGANISM: synthetic construct
US-09-822-250-14
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RESULT 19
US-10-044-674-86/c
US-10-044-674-86/c
Sequence 86, Application US/10044674
Publication No. US20030175710A1
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; Sequence 1647, Application US/10033145
; Publication No. US20020151515A1
; GENERAL INFORMATION:
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Best Local Similarity
""" | Gonserv
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Best Local Similarity
Watches 9; Conserva
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 1647
LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES FILE REFERENCE: GAOZOIC CURRENT APPLICATION NUMBER: US/10/033,145
PRIOR APPLICATION 2001-11-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: PCT/US99/13800 PRIOR FILING DATE: 1999-06-18
                                                                                                                                                                                                                                                                  TYPE: DNA
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100.0%; Pred. No.
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SEQ ID NO 86
LENGTH: 10
                                                                                                                                       REGISTRATION NUMBER: P44,170
REFERENCE/DOCKET NUMBER: CXU-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (864) 271-1592
TELEFAX: (864) 233-7342
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 6, Application US/09758073
Patent No. US20010024785A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Nandabalan, Krishnan
APPLICANT: Stephens, J. Claiborne
TITLE OF INVENTION: HAPLOTYPES OF THE TNFRSF11B GENE
FILE REFERENCE: TNFRSF11B MWH-0001US (CIP)
CURRENT APPLICATION NUMBER: US/10/044,674
CURRENT FILING DATE: 2002-01-09
PRIOR APPLICATION NUMBER: PCT/US00/18803
PRIOR APPLICATION STEP PCT/US00/18803
PRIOR APPLICATION NUMBER: PCT/US00/18803
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 94
                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM compatible
OPERATING SYSTEM: MS Dos; W
SOFTWARE: WordDerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/7
FILING DATE: Filed Herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Chew, Anne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Keinath, et al.
TITLE OF INVENTION: Method of Diagnosing Gummy Stem Blight in
TITLE OF INVENTION: Plants Using a Polymerase Chain Reaction Assay
                                                                                                                                                                                                                                                                                  FILING DATE: 16-MAR-1998
ATTORNEY/AGENT INFORMATION:
NAME: Judy C. Jarecki-Black, Ph.D.
                                    TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
MOLECULE TYPE: Other Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
DESCRIPTION: HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Judy C. Jarecki-Black, Ph.D.
ADDRESSEE: Dority & Manning, P.A.
STREET: 700 E. No. US20010024785Alth Street,
CITY: Greenville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                   ENGTH:
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                   Oligonucleotide Primer
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100.0%; Pr
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Query Match
Best Local Similarity
Watches 9; Conserve
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Best Local Similarity
Tatches 9; Conserve
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; OTHER INFORMATION: Exon/intron of TORB
US-09-772-105-77
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US-09-772-105-77/c
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                                                                                                                                                                                                                                                                                         US-10-033-145-1651
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: F
SEQ ID NO 77
LENGTH: 10
                                                                                                                                                                                     Sequence 1651, Application US/10033145
Publication No. US20020151515A1
GENERAL INFORMATION:
APPLICANT: GENZYME CORPORATION
APPLICANT: ROBERTS, BRUCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION
NUMBER OF SEQ ID NOS: 2137
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1651
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Patent No. US20010029015A1
                                                   CURRENT APPLICATION NUMBER: US/10/033,145
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
                                                                                                                               APPLICANT: SHANKARA, STRINIVAS
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
FILE REFERENCE: GA0201C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 1997-06-19 NUMBER OF SEQ ID NOS: 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/218,363
PRIOR FILING DATE: 1998-12-22
PRIOR APPLICATION NUMBER: US 09/099,454
PRIOR FILING DATE: 1998-06-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/772,105
CURRENT FILING DATE: 2001-01-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Ozelius, Laurie J.
APPLICANT: Breakefield, Xandra O.
TITLE OF INVENTION: TORSIN, TORSIN-RELATED GENES, AND
TITLE OF INVENTION: METHODS OF DETECTING NEURONAL DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
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Operon Technologies
(B: No. US20010024785A1 Applicable
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Pred. No. 17;
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PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 292
LENGTH: 10
TYPE: DNA
OPGRANTON
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US-10-330-627-292
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US-10-330-627-292
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APPLICANT: Kinzler, Kenneth W
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
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Best Local S
Matches 9
                                                   Query Match 28.0%;
Best Local Similarity 90.0%;
Matches 9; Conservative
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APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
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ORGANISM: Homo sapiens
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1499 TCAGCAGCCA 1508
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b. US20030175771A1
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p. US20030175771A1
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90.0%;
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                                                                       Score 8.4; DB 1; Length 10; Pred. No. 17;
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Pred. No. 17;
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Pred. No. 1
                                                     Mismatches
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US-10-330-627-1077/c
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Best Local Similarity 90.0
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1077
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CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
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TITLE OF INVENTION: Human Tra
FILE REFERENCE: 001107.00319
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                                                                                                       APPLICATION NUMBER: US/08/894,454
FILING DATE: 15-AUG-1997
APPLICATION NUMBER: PCT/EP/00716
FILING DATE: 19-FEB-1996
APPLICATION NUMBER: 95200390.3
FILING DATE: 17-FEB-1995
APPLICATION NUMBER: 95201951.1
FILING DATE: 14-JUL-1995
APPLICATION NUMBER: 95201951.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: OSS
COMPUT
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CORRESPONDENCE ADDRESS:
ADDRESSEE: The Webb Law Firm
ADDRESSEE: The Webb Law Firm
ADDRESSEE: The Webb Law Firm
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CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA: US/10/352,615
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TITLE OF INVENTION: MULTIPLE-TUMOR ABERRENT GROWTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1490 AGCCAGACTT 1499
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REFERENCE/DOCKET NUMBER: 702-971100
                                  NAME: Johnson, Barbara E
REGISTRATION NUMBER: 31,1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ
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90.0%;
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Pred. No.
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RESULT 27
US-09-862-847-15/c
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.1
SEQ ID NO 15
LENGTH: 11
TYPE: DNA
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                                APPLICANT: Webb, Suzin M.

APPLICANT: Affourtit, Jason P.

TITLE OF INVENTION: GENETIC POLYMORPHISMS IN THE HUMAN NEUROKININ 1 RECEPTOR GENE AND TITLE OF INVENTION: USES IN DIAGNOSIS AND TREATMENT OF DISEASES

FILE REFERENCE: PC10461AGPR
CURRENT APPLICATION NUMBER: US/10/146,354A

CURRENT FILING DATE: 2002-08-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WILLING DATE: 2001-04-20 NUMBER OF SEQ ID NOS: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/862,847
CURRENT FILING DATE: 2001-05-21
PRIOR APPLICATION NUMBER: US 60/206,537
PRIOR FILING DATE: 2000-05-21
                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                    APPLICANT: Pfizer Inc
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INFORMATION FOR SEQ ID NO: 110:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Synthetic oligonucleotide primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
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APPLICATION NUMBER: 60/293,425 FILING DATE: 2001-05-25
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                              Seymour, Albert B.
Nelson, Darcy L.
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Pred. No. 17;
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Query Match
Best Local Similarity
Matches 10; Conserv:
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                                                           US-09-989-789-455
                                                                                                                                                                                                                                                                                                                                                US-09-989-789-455/c
                                                                                                                                             CURRENT APPLICATION NUMBER: US/03
CURRENT FILING DATE: 2002-03-25
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 455
LENGTH: 9
                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-C011.20 / S11-US2
                                                                                                                                                                                                                                                                                                                 Sequence 455, Application US/09989789 Patent No. US20020063379A1
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LENGTH: 20
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Publication No.
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            Best Local Similarity
                             Query Match
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CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 FILE REFERENCE: RTS-0250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Brett P. Monia
APPLICANT: Susan M. Frei
APPLICANT: Scott Cooper
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ORGANISM: Homo sapiens
                                                                         OTHER INFORMATION: Description of Artificial Sequence: example target OTHER INFORMATION: DNA
                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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                                                                                                      FEATURE:
                                                                                                                                     TYPE: DNA
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Scott Cooper
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76.9%;
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              Score 8; 1; Pred. No.
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Pred. No. 29,
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Pred. No. :
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1493 CAGACTTC 1500

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US-09-990-186-456/c
; Sequence 456, Application US/09990186
; Publication No. US20030068675A1
; GENERAL INFORMATION:
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US-09-990-186-455/c
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                                                                             RESULT 33
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US-09-989-789-456/c
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; Publication No. US200
; GENERAL INFORMATION:
; APPLICANT: LIU, Qian
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TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/09/989,789
CURRENT FILING DATE: 2002-03-25
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 456
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Matches
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Best Local Similarity
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                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                             -09-990-186-455
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APPLICANT: LIU, Qiang
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TYPE: DNA
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TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
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o. US20030068675A1
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100.0%; Pr
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Sequence 456, Application US/09989994

Publication No. US20030104526A1

GENERAL INFORMATION:
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.0/ S11.US2
CURRENT APPLICATION NUMBER: US/09/989,994

CURRENT FILING DATE: 2001-11-20

NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver: 2.0
SEQ ID NO 456
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; OTHER INFORMATION:
US-09-990-186-456
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US-09-989-994-456/c
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Best Local S
Matches 8
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TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.21 / $11-US3
CURRENT APPLICATION NUMBER: US/09/990,186
CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 456
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LENGTH: 9
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Best Local
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TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/09/989,994
CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
COFTWARE: PETENTIN Ver. 2.0
                  TYPE: DI
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ORGANISM: Artificial Sequence
FEATURE:
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ORGANISM: Artificial Sequence
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100.0%; Pred. No.
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Query Match
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               SEQ ID NO 6
LENGTH: 9
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Matches 8; Conserv
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Publication No. US20030162211A1
                                                                        TITLE OF INVENTION: METHOD AND KIT FOR THE DETERMINATION OF TITLE OF INVENTION: CELLULAR ACTIVATION PROFILES FILE REFERENCE: VANM212.001CP1
CURRENT APPLICATION NUMBER: US/10/339,161
CURRENT FILING DATE: 2003-01-07
PRIOR APPLICATION NUMBER: US 09/816,763
PRIOR APPLICATION NUMBER: US 09/816,763
PRIOR FILING DATE: 2001-03-23
PRIOR FILING DATE: 2001-03-23
PRIOR FILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Fang, Yu
APPLICANT: Wang, Xiao-Yang
APPLICANT: Wang, Xiao-Yang
APPLICANT: Turpin, Pierre
TITLE OF INVENTION: Methods of detecting multiple DNA
TITLE OF INVENTION: binding protein and DNA interactions in a sample,
TITLE OF INVENTION: devices, systems and kits for practicing the same.
FILE REFERENCE: CLON-071
CURRENT APPLICATION NUMBER: US/10/113,877
CURRENT FILING DATE: 2002-03-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/314,330
PRIOR FILING DATE: 2001-08-20
NUMBER OF SEQ ID NOS: 192
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                           SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                          APPLICANT: Remacle, Jose
APPLICANT: Renard, Patricia
APPLICANT: Art, Muriel
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TYPE: DNA
ORGANISM: Artificial Sequence
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TYPE: DNA
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100.0%; Pred. No. 95
tive 0; Mismatches
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TOTHER INFORMATION: Transcription factor SMAD 4
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US-10-277-494-147/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or conditions Related
TITLE OF INVENTION: Epidermal Growth Factor Receptors
TITLE OF INVENTION: Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-K (400/064)
CURRENT FILING DATE: 2002-10-21
CURRENT FILING DATE: 2002-10-21
NUMBER OF SEQ ID NOS: 446
SOFTWARE: PatentIn version 3.0
SEQ ID NO 147
LENGTH: 9
TYPE: RNA
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Matches 8; Conserv
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                                 CURRENT APPLICATION DATA:
                                                                                                   ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Ly
                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                 COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                       STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
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nes 8; Conserv
                APPLICATION NUMBER:
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milarity 100.0%;
Conservative 0
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Woodring E. Wright
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MBER: US/08/463,404
05-JUN-1995
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Pred. No. 95;
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Levels

PRIOR APPLICATION DATA:

08/060,952

CLASSIFICATION: APPLICATION NUMBER:

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GENERAL INFORMATION:
APPLICANT: GENZYME CORPORATION
APPLICANT: ROBERTS, BRUCE
APPLICANT: SHANKARA, SRINIVAS
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
FILE REFERENCE: GAO201C
CURRENT APPLICATION NUMBER: US/10/033,145
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
NUMBER OF SEQ ID NOS: 2137
SOFTMARE: PATENTIN VERSION 3.0
SEQ ID NO 185
LENGTH: 10
TYPE: DNA
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                                                                                                                                RESULT 41
US-10-033-145-1011
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US-10-033-145-185
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                                                                         Sequence 1011, Application US/10033145
Publication No. US20020151515A1
GENERAL INFORMATION:
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Best Local (
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                     APPLICANT: GENZYME CORPORATION APPLICANT: ROBERTS, BRUCE APPLICANT: SHANKARA, SRINIVAS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 202/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
PPLICANT: SHANKARÀ, SRINIVAS
ITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: May 13, 1993
APPLICATION NUMBER: 07/882,438
FILING DATE: May 13, 1992
APPLICATION NUMBER: 08/038,766
FILING DATE: March 24, 1993
ATTORNEY/AGENT INFORMATION:
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Local Similarity 100.0%; I
nes 8; Conservative 0;
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Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: DNA Template US-10-113-030-3
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Sequence 3, Application US/10358818
Publication No. US20030162213A1
GENERAL INFORMATION:
APPLICANT: Fuller, Carl
APPLICANT: Kulmar, Shiv
APPLICANT: Sood, Anup
APPLICANT: Sood, Anup
APPLICANT: Nelson, John
TITLE OF INVENTION: Terminal-Phosphate-Labeled Nucleotides and Methods of Use
FILE REFERENCE: PB0156-1CIP
CURRENT APPLICATION NUMBER: US/10/358,818
CURRENT FILING DATE: 2003-02-05
PRIOR APPLICATION NUMBER: US 60/315,798
PRIOR APPLICATION NUMBER: US 60/315,798
PRIOR APPLICATION NUMBER: US 10/113,030
PRIOR FILING DATE: 2001-08-29
PRIOR APPLICATION NUMBER: US 10/113,030
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APPLICANT: Sood, Anup
APPLICANT: Sood, Sood, Anup
APPLICANT: Sood, Anup
APPLICANT: Kumar, Shiv
TITLE OF INVENTION: Terminal-Phosphate-Labeled Nucleotides and Methods of Use
FILE REFERENCE: PB0156-1
CURRENT APPLICATION NUMBER: US/10/113,030
CURRENT FILING DATE: 2002-04-01
PRIOR APPLICATION NUMBER: US 60/315,798
PRIOR APPLICATION NUMBER: US 60/315,798
PRIOR TILING DATE: 2001-08-29
NUMBER OF SEQ ID NOS: 3
SOFTWARE: Patentin version 3.1
SEQ ID NO 3
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Publication No. US20030077610A1
GENERAL INFORMATION:
APPLICANT: Nelson, John
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Best Local Similarity lov.
Thickes 8; Conservative
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Matches 8; Conserv
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CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
NUMBER OF SEQ ID NOS: 2137
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100.0%; Pred. No.
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Sequence 344, Application US/10330627

Publication No. US20030175771A1

GENERAL INFORMATION:

APPLICANT: Velculescu, Victor E.

APPLICANT: Velculescu, Victor E.

APPLICANT: Vegelstein, Bert

TITLE OF INVENTION: Human Transcriptomes

FILE REFERENCE: 001107.00319

CURRENT APPLICATION NUMBER: US/10/330,627

CURRENT APPLICATION NUMBER: US 09/448,480

PRIOR APPLICATION NUMBER: US 09/448,480

PRIOR FILING DATE: 1999-11-24

NUMBER OF SEQ ID NOS: 1564

SOFTMARE: FastSEQ for Windows Version 4.0

SEQ ID NO 344

LENGTH: 10

TYPE: DNA

ORGANISM: Homo Sapiens

US-10-330-627-344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; NUMBER OF SEQ ID NOS: 3
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 3
; LENGTH: 10
TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DNA Template
US-10-358-818-3
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US-10-330-627-344/c
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Search completed: April 15, 2004, 16:38:43 Job time: 0.001 secs
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RESULT 1
BG896271
LOCUS
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AUTHORS
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ORGANISM
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KEYWORDS
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Maximum Match 100%
Listing first 1 summaries
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                                                                               COMMENT
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GlaxoSmithKline
709 Swedeland Road, P.O.
Tel: 610-270-7245
                                                                                                                                  Identification and initial characterization of 5000 expressed sequenced tags (ESTs) each from adult human normal and osteoarthritic cartilage cDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (2001)
                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 11)

Kumar, S., Connor, J.R., Dodds, R.A., Halsey, W., Van Horn, M., Mao, J., Sathe, G., Mui, P., Agarwal, P., Badger, A.M., Lee, J.C., Gowen, M. and Lark, M.W.
                                                           UW2109
                                                                           Contact: Sanjay Kumar
                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                            BG896271.1 GI:14306512
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Email: sanjay_kumar-l@gsk.com
Seq primer: T7.
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                                                                                                                  Conservative
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/lab_host="E.coli DH10 B"
/clone_lib="HOA (Human Osteoarthritic Cartilage)"
/note="Vector: pSPORT I; Site_1: SalI; Site_2: NotI;
Directional"
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/db_xref="taxon:9606"
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